# **IDWeek 2022** Poster #P2044

# Michael D. Huband<sup>1</sup>, Michael A. Pfaller<sup>1,2</sup>, Jennifer M. Streit<sup>1</sup>, Helio S. Sader<sup>1</sup>, and Mariana Castanheira<sup>1</sup> <sup>1</sup> JMI Laboratories, North Liberty, Iowa, United States; <sup>2</sup> University of Iowa, Iowa City, Iowa, United States RESULTS Omadacycline and comparator agent MIC and susceptibility data against key Gram-

## INTRODUCTION

- Omadacycline is a novel aminomethylcycline tetracycline antibacterial with oral and intravenous formulations that was approved by the United States Food and Drug Administration (FDA) for treatment of adults with acute bacterial skin and skin structure infections (ABSSSI) and community-acquired bacterial pneumonia (CABP) caused by indicated organisms.
- Ongoing clinical trials for omadacycline include for the treatment of diabetic foot infection (NCT04714411), community-acquired bacterial pneumonia (NCT04779242), and pulmonary disease caused by *M. abscessus* complex (NCT04922554) as well as for the evaluation of pharmacokinetics in cystic fibrosis (NCT04460586) and pharmacokinetics of intravenous and oral omadacycline in children and adolescents with suspected or confirmed bacterial infections (NCT05217537).
- Omadacycline has previously demonstrated potent in vitro activity against Grampositive (staphylococci, streptococci, and enterococci) and Gram-negative (Enterobacter cloacae, Haemophilus influenzae, Klebsiella pneumoniae, and Escherichia coli) bacterial pathogens commonly associated with ABSSSI, communityacquired respiratory tract infection (CARTI), and pneumonia in hospitalized patients (PIHP), including isolates expressing common tetracycline-, penicillin/oxacillin-, fluoroquinolone-, and macrolide-resistance mechanisms.
- The activity of omadacycline and tetracycline comparators against 14,000 recent (2020– 2021) bacterial isolates collected from patients in United States medical centers (SENTRY Antimicrobial Surveillance Program) and stratified by infection type is presented.

# MATERIALS AND METHODS

- 14,000 bacterial isolates were recovered (1 per patient infection episode) from patients with documented infections in 31 United States medical centers in 9 Census Divisions representing multiple infection types.
- The isolates included 3,769 Staphylococcus spp., 1,231 Streptococcus spp., 720 Enterococcus spp., 466 Haemophilus spp., 211 Moraxella spp., 1,830 non-fermenters, and 5,773 Enterobacterales.
- Isolates were collected from patients with skin and skin structure infections (SSSI; 3,131 isolates; 22.4%), bloodstream infections (BSI; 3,371 isolates; 24.1%), CARTI (1,162 isolates; 8.3%), intra-abdominal infections (IAI; 776 isolates; 5.5%), PIHP (3,167 isolates; 22.6%), urinary tract infections (UTI; 2,224 isolates; 15.9%), and other infections (169 isolates; 1.2%).
- Bacterial identifications were confirmed at JMI Laboratories using standard microbiology methods and matrix-assisted laser desorption ionization-time of flight mass spectrometry (Bruker Daltonics, Bremen, Germany).
- Broth microdilution susceptibility testing was performed according to Clinical and Laboratory Standards Institute M07 (CLSI; 2018) methodology.
- Susceptibility results were interpreted using CLSI M100 (2022), European Committee on Antimicrobial Susceptibility Testing (EUCAST; v12.0, 2022), and FDA breakpoint interpretive criteria.

## Table 1. Antimicrobial activity of omadacycline and tetracycline comparators against bacterial isolates collected from patients with skin and skin structure infections in United States medical centers (2020–2021)

		Omadacycline			Tetracycline		Tigecycline		Do	xycycline		Omadacycline		Tetra	Tetracycline		Tigecycline		Doxycycline			Omadacycline		Tetra	Tetracycline		Tigecycline		Doxycycl	
Organism (no. tested)	MIC	50 / 90	%S (FDA)	MI	MIC <sub>50 / 90</sub> %S		MIC <sub>50 / 5</sub>	● %S (FDA/ ■ EUCAST)	Α/ Γ) MIC <sub>50/9</sub>	<ul><li>%S (CLSI/</li><li>EUCAST)</li></ul>	Organism (no. tested)		50 / 90			%S (CLSI/ EUCAST)					∕₀S (CLSI/	Organism (no. tested)	-	• (FDA				%S (FDA/ EUCAST)		0
Staphylococcus aureus (1,575)	0.12	0.12	99.0	≤0.5	1	92.1 / 90.2	0.06 (	.12 100.0 / 10	0.0 ≤0.06 (	0.5 97.1 / 95.1			50 / 90	(FDA)	50 / 90	EUCAST)	50 / 90	EUCAST)	50	/ 90	EUCAST)		50 / 90	(FDA	50 / 90	EUCAST)	50 / 90	EUCAST)	50 / 90	
MRSA (659)	0.12	0.25	98.0	≤0.5	8	88.0 / 86.5	0.06 0	.12 100.0 / 10	0.0 ≤0.06	1 95.1 / 92.4	Streptococcus pneumoniae (594)	0.06	0.06	99.8	0.25 >4	79.4 / 79.4	0.03 0.06	98.3 / —ª	0.12	>1 7	78.6 / 80.5	Staphylococcus aureus (832)	0.12 0.2	<b>25</b> 94.4	≤0.5 ≤0.5	94.7 / 92.4	0.06 0.12	100.0 / 100.0	) ≤0.06 0.5	5 §
MSSA (916)	0.12	0.12	99.8	≤0.5	≤0.5	95.1 / 92.6	0.06 0	.12 100.0 / 10	0.0 ≤0.06 0	.12 98.5 / 97.0																				
S. lugdunensis (41)	0.06	0.06	97.6	≤0.5	≤0.5	95.1 / 95.1	0.03 (	.06 — <sup>b</sup> / 100	.0 ≤0.06 ≤0	0.06 100.0 / 95.1	S. pneumoniae penicillin-R <sup>b</sup> (64)	0.06	0.06	100.0	0.25 >4	51.6 / 51.6	0.06 0.06	98.4 / —ª	0.25	>1 5	50.0 / 53.1	MRSA (321)	0.12 <b>0</b> .	<b>.5</b> 86.0°	<sup>a</sup> ≤0.5 2	90.9 / 87.8	0.06 0.25	100.0 / 100.0	/ ≤0.06 1	Q
Streptococcus anginosus group (22)ª	0.06	0.06	95.5	4	>4	40.9 / <u></u>	0.03 (	.06 100.0 / -	_ <sup>b</sup> 0.5	>1b /b	S. pneumoniae macrolide-R (270)	0.06	0.06	100.0	0.25 >4	58.4 / 58.4	0.06 0.06	98.9 / <u></u> a	0.12	>1 5	57.4 / 59.6	MSSA (511)	0.12 0.1	12 99.6	≤0.5 ≤0.5	97.1 / 95.3	0.06 0.12	100.0 / 100.0	) ≤0.06 0.1′	2 9
S. pyogenes (104)	0.06	0.12	100.0	0.25	>4	66.3 / 66.3	0.06 0	.06 100.0 / 10	0.0 0.12	>1b / 66.3																				
S. pyogenes macrolide-R (28)	0.06	0.12	100.0	>4	>4	7.1 / 7.1	0.06 0	.06 100.0 / 10	0.0 >1 :	>1b / 7.1	S. pneumoniae tetracycline-R (119)	0.06	0.06	100.0	>4 >4	0.0 / 0.0	0.06 0.06	98.3 / —ª	>1	>1	0.8/2.5	Streptococcus pneumoniae (12)	0.06 0.0	<b>J6</b> 100.0	0.25 >4	83.3 / 83.3	0.06 0.06	100.0 / <u> </u> °	0.12 >1	8
S. pyogenes tetracycline-R (35)	0.06	0.12	100.0	>4	>4	0.0 / 0.0	0.06	.06 100.0 / 10	0.0 >1 :	>1b / 0.0	Haemophilus influenzae (371)	0.5	1	99.7	0.5 0.5	98.9 / 98.9	0.25 0.25	90.8 / <u> </u>			a /a	Haemophilus influenzae (32)	0.5 1	100./	0.5 0.5	100.0 / 100.0	0.12 0.25	100.0		_
Enterococcus faecalis (114)	0.06	0.12	100.0	>16	>16	26.3 / — <sup>b</sup>	0.06	.12 100.0 / 10	0.0 8	8 49.1 / — <sup>b</sup>																				
<i>E. faecalis</i> vancomycin-R (4)	0.12	—	100.0	>16		0.0 / <u> </u> b	0.06	— 100.0 / 10	0.0 8	8 0.0 / — <sup>b</sup>	Moraxella catarrhalis (193)	≤0.12	0.25	100.0 <sup>c</sup>	0.25 0.5	99.0 / 99.0	0.06 0.06	<u> </u>			<u>     a  /      a</u>	A. baumannii-calcoaceticus species	0.5 4	<b>4</b> 90.0°	2 >16	67.3 / <u></u>	0.25 4	98.2° / — <sup>b</sup>	0.12 >8	3
E. faecium (30)	0.06	0.12	b	>16	>16	6.7 / — <sup>b</sup>	0.06	.12 — <sup>b</sup> / 100	.0 8	8 40.0 / — <sup>b</sup>	R, resistant; S, susceptible. <b>Bold</b> , omadacycline MIC <sub>90</sub> values.											complex (110)								
<i>E. faecium</i> vancomycin-R (21)	0.06	0.12	b	>16	>16	9.5 / — <sup>b</sup>	0.06	.12 — <sup>b</sup> / 100	.0 8	8 38.1 / — <sup>b</sup>		Green, susceptible according to CLSI or FDA breakpoint interpretive criteria.   Green, susceptible according to CLSI or FDA breakpoint interpretive criteria.   Klebsiella pneumoniae (281) 2 8.3 1 >16 77.2 / -b 0.5 1 96.8 / -b											2 >8	3						
Enterobacter cloacae species complex <sup>d</sup> (88)	2	8	87.5	2	16	81.8 / — <sup>b</sup>		1 96.6 / —		>8 80.7 / — <sup>b</sup>	<sup>a</sup> Breakpoint interpretive criteria were unavailable.	<sup>a</sup> Breakpoint interpretive criteria were unavailable.																		
Klebsiella pneumoniae (99)	2	8	88.9	2	>16	70.7 / — <sup>b</sup>	0.5	1 99.0 / —	- <sup>b</sup> 2 >	>8 71.7 / — <sup>b</sup>	<sup>b</sup> Oral dosing (penicillin MIC, ≥2 mg/L). <sup>c</sup> % inhibited at ≤0.25 mg/L.											Escherichia coli (214)	0.5 2	<u>/</u> 99.1°	2 >16	59.8 / <u></u>	0.12 0.25	100.0 / 99.5	1 >8	, (
Escherichia coli (174)	0.5	2	97.1 <sup>c</sup>	2	>16	62.6 / <u></u>	0.12 (	.25 100.0 / 10	0.0 1	>8 68.4 / <u>b</u>												MRSA, methicillin-resistant S. aureus; MSSA, methicillin-sus	ceptible S. aureus; S	З, susceptible.						
MRSA, methicillin-resistant <i>S. aureus</i> ; MSSA, methicillin-susceptible <i>S. au</i> <b>Bold</b> omadacycline MIC <sub>60</sub> values	<i>ureus</i> ; R, resista	ant; S, suscep	tible																			<b>Bold</b> , omadacycline MIC <sub>90</sub> values. Green, susceptible according to CLSI or FDA breakpoint inte	rpretive criteria.							

**Bold**, omadacycline  $MIC_{90}$  values. Green, susceptible according to CLSI or FDA breakpoint interpretive criteria

ellow, intermediate according to CLSI or FDA breakpoint interpretive criteria Gray, resistant according to CLSI, EUCAST, or FDA breakpoint interpretive criteria.

Includes S. anginosus group (20) and S. constellatus (2). <sup>b</sup> Breakpoint interpretive criteria were unavailable.

° % inhibited at ≤2 mg/L.

<sup>d</sup> Includes *E. cloacae* (30) and *E. cloacae* species complex (58)

- positive and Gram-negative bacterial pathogens collected from patients with SSSI, CARTI, and PIHP in United States medical centers during 2020–2021 are presented in Tables 1–3.

- from PIHP (Table 1).

- (Table 1).
- (Tables 2–3).
- CARTI (Table 2).

- (Table 3

Bacterial pathogen occurrence by infection type for omadacycline is presented in Figures 1–3 for SSSI, CARTI, and PIHP, respectively.

Omadacycline demonstrated potent *in vitro* activity against methicillin-resistant S. aureus (MRSA) and methicillin-susceptible S. aureus (MSSA) from SSSI (MIC<sub>50/90</sub> values of 0.12/0.12-0.25 mg/L) with percent susceptibility (%S) of 98.0%S (FDA) and 99.8%S (FDA), respectively (Table 1).

- Omadacycline (MIC<sub>50/90</sub>, 0.12/0.12 mg/L) was active against 99.6% of MSSA isolates

97.6% of S. lugdunensis isolates from SSSI (MIC<sub>50/90</sub>, 0.06/0.06 mg/L) were susceptible to omadacycline (FDA; Table 1).

Omadacycline (MIC<sub>50/90</sub>, 0.06/0.06 mg/L; 95.5%S [FDA]) and tigecycline (MIC<sub>50/90</sub>, 0.03/0.06 mg/L; 100.0%S [EUCAST]) demonstrated activity ≥90.0% against S. anginosus group isolates from SSSI (Table 1).

All S. pyogenes isolates from SSSI were susceptible to omadacycline (MIC<sub>50/90</sub>, 0.06/0.12 mg/L; 100.0%S [FDA]), including macrolide- and tetracycline-resistant strains

All *E. faecalis* isolates from SSSI were susceptible to omadacycline (MIC<sub>50/90</sub>, 0.06/0.12 mg/L; 100%S [FDA]), including vancomycin-resistant strains (Table 1). Omadacycline was active against S. pneumoniae isolates from CARTI (MIC<sub>50/00</sub>, 0.06/0.06 mg/L; 99.8%S [FDA]) and PIHP (MIC<sub>50/90</sub>, 0.06/0.06 mg/L; 100%S [FDA])

- Omadacycline (MIC<sub>50/90</sub>, 0.06/0.06 mg/L) was active against 100% of penicillinresistant, macrolide-resistant, and tetracycline-resistant S. pneumoniae isolates from

Omadacycline was active against *H. influenzae* isolates from CARTI (MIC<sub>50/90</sub>, 0.5/1 mg/L; 99.7%S [FDA]) and PIHP (MIC<sub>50/90</sub>, 0.5/1 mg/L; 100%S [FDA]) (Tables 2–3). Susceptibility of *E. cloacae* species complex and *K. pneumoniae* isolates from SSSI to omadacycline was 87.5%S and 88.9%S, respectively (Table 1).

- 88.3% of *K. pneumoniae* isolates from PIHP were susceptible to omadacycline

97.1% of *E. coli* from SSSI and 99.1% of *E. coli* from PIHP were inhibited by  $\leq 4 \mu g/mL$ of omadacycline (Tables 1 and 3).

90.0% of *A. baumannii-calcoaceticus* species complex isolates from PIHP were inhibited by  $\leq 4 \mu g/mL$  of omadacycline (Table 3).

(*n*=1,162; 2020–2021)

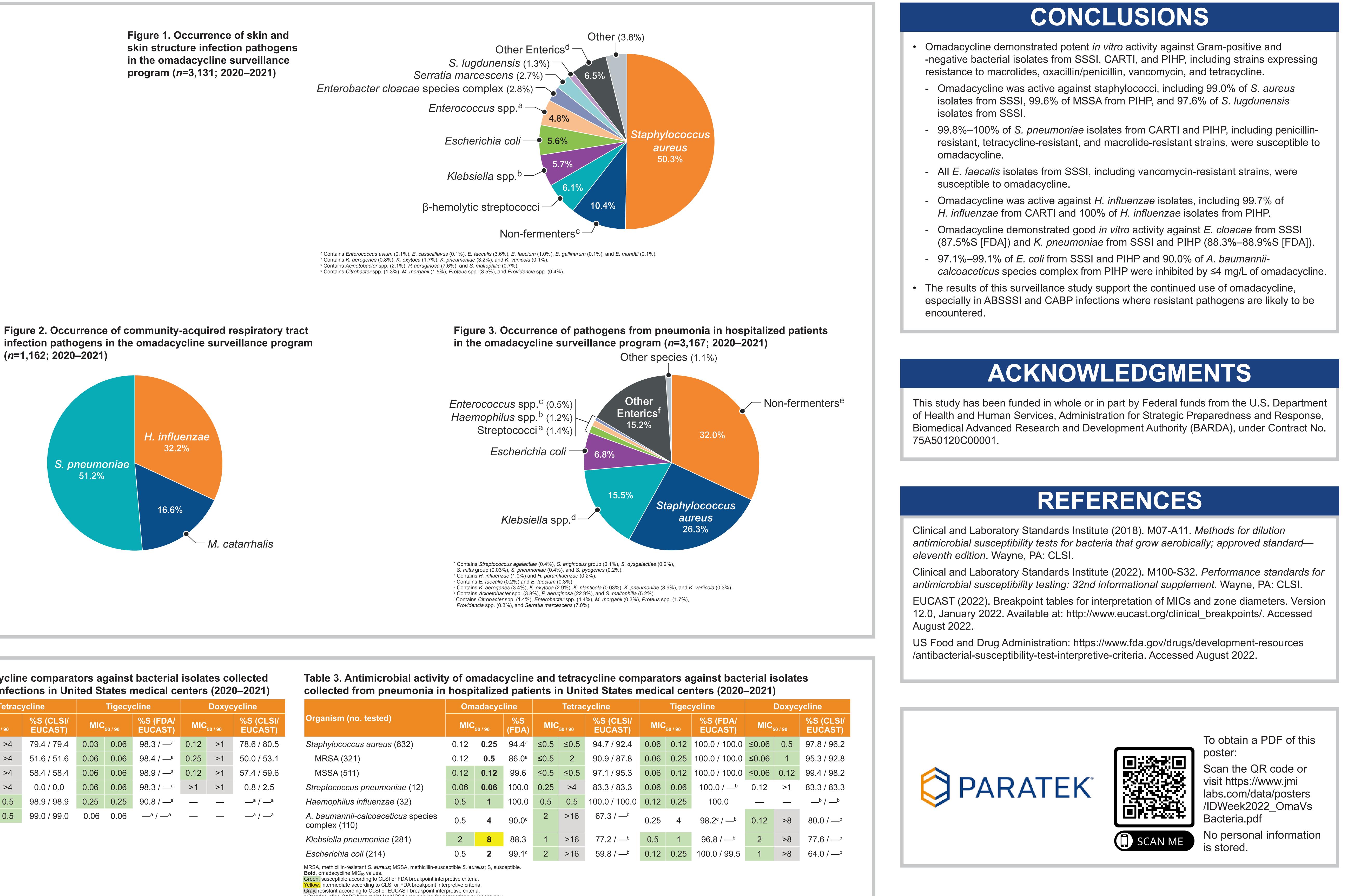
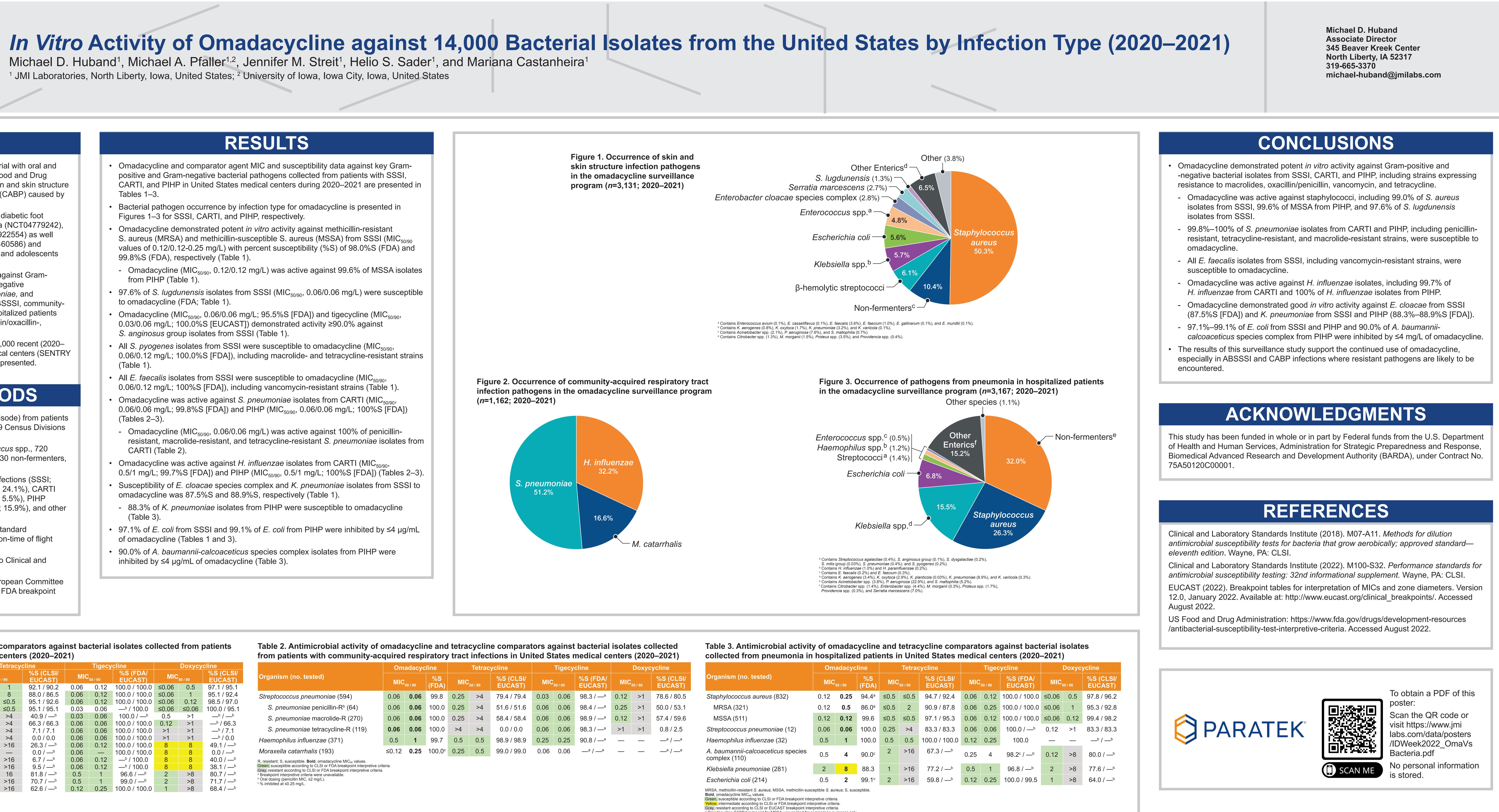


Table 2. Antimicrobial activity of omadacycline and tetracycline comparators against bacterial isolates collected from patients with community-acquired respiratory tract infections in United States medical centers (2020–2021)



Omadacycline CABP breakpoint for MSSA was applied for comparison purposes only

<sup>b</sup> Breakpoint interpretive criteria were unavailable  $^{\circ}$  % inhibited at  $\leq 4$  mg/L.