

# Microbiology and Preclinical Review of Omadacycline

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Omadacycline is a novel aminomethylcycline antimicrobial and semisynthetic derivative of tetracycline. In vitro, omadacycline displays potent activity against gram-positive and many gram-negative bacteria, including methicillin-resistant *Staphylococcus aureus*, *Streptococcus pneumoniae*,  $\beta$ -hemolytic streptococci, vancomycin-resistant *Enterococcus*, and Enterobacteriaceae. Omadacycline is also active against atypical and anaerobic pathogens, including *Legionella pneumophila*, *Mycoplasma* spp., *Ureaplasma* spp., *Bacteroides* spp., and *Clostridioides difficile*. This review outlines the microbiology and preclinical studies of omadacycline, including its mechanism of action; spectrum of activity; protein binding; activity in the presence of surfactant, serum, normal, and pH-adjusted urine, or bacterial biofilms; postantibiotic effect; pharmacodynamic properties; and in vitro and in vivo efficacy. The results of in vitro and in vivo animal studies support the observations made in phase III clinical trials and the clinical development of omadacycline.

**Keywords.** antimicrobial; omadacycline; pharmacodynamics; spectrum of activity; tetracyclines.

Omadacycline is an aminomethylcycline antimicrobial of the tetracycline class, approved in the United States for the treatment of community-acquired bacterial pneumonia and acute bacterial skin and skin structure infections (ABSSSI) in adults [1]. Omadacycline has demonstrated in vitro activity against resistant gram-positive pathogens, including methicillin-resistant *Staphylococcus aureus* (MRSA), penicillin- and macrolide-resistant *Streptococcus pneumoniae*, vancomycin-resistant *Enterococcus* (VRE), and *Clostridioides difficile*, pathogens that have been recognized by the US Centers for Disease Control and Prevention as urgent or serious threats [2]. The efficacy of omadacycline as monotherapy for serious community-acquired bacterial infections, including ABSSSI and community-acquired bacterial pneumonia, has been demonstrated in phase III clinical trials [3–5]. Ongoing clinical trials are investigating omadacycline for the treatment of urinary tract infections.

## MECHANISM OF ACTION

Omadacycline is a semisynthetic tetracycline derivative and displays the same mechanism of action as the tetracycline class. Like tetracyclines, omadacycline inhibits bacterial protein synthesis by binding the primary tetracycline binding site on the 30S subunit of the bacterial ribosome [6]. As demonstrated

by whole-cell, macromolecular synthesis, omadacycline directly inhibits bacterial protein synthesis while sparing bacterial DNA, RNA, and peptidoglycan synthesis [7]. Through modifications at the C-7 and C-9 positions of the tetracycline D-ring, omadacycline is able to overcome common tetracycline resistance mechanisms, including tetracycline-specific efflux pumps and ribosomal protection (Figure 1). The C-7 modification circumvents the tetracycline-specific efflux pump resistance mechanism, whereas the C-9 modification overcomes the ribosomal protection resistance mechanism. Although the binding of omadacycline to the primary bacterial ribosome site is similar to that of other tetracyclines, dimethyl sulfate chemical probing and Fenton cleavage studies have revealed that omadacycline, tetracycline, and tigecycline have unique, nonspecific interactions with the 16S rRNA [6]. The unique interaction of omadacycline with the bacterial ribosome may help to explain its ability to overcome the standard tetracycline resistance mechanisms. Omadacycline retains activity against gram-positive pathogens that carry resistance genes for ribosomal protection (ie, *tetM*, *tetO*, and *tetS*) and tetracycline efflux (ie, *tetK* and *tetL*) [7, 8]. Thus, omadacycline remains active against tetracycline-resistant bacterial strains. To date, the resistance mechanisms that have been found to inhibit the activity of omadacycline include multidrug efflux pumps (MexXY-OprM and MexAB-OprM) [9] and the tetracycline monooxygenase TetX (Paratek Pharmaceuticals, Inc.; data on file). Although TetX has been shown to inactivate all known tetracyclines, it is not a widespread resistance determinant [10, 11].

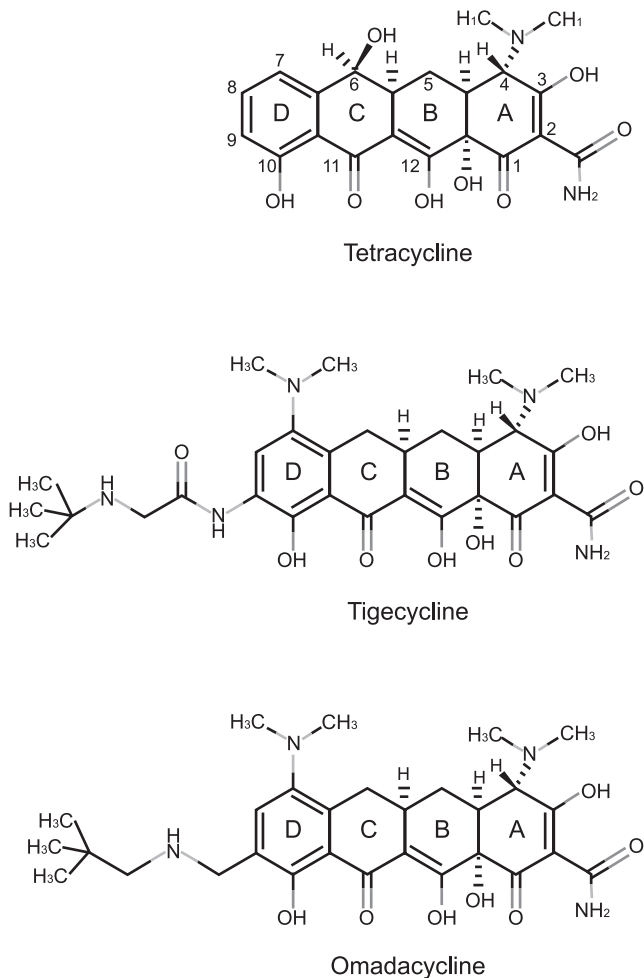
## SPECTRUM OF ACTIVITY

Omadacycline has been evaluated in a range of studies, including centralized US clinical isolate surveillance studies that began in 2010. Consistency in the spectrum of omadacycline

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**Figure 1.** Chemical structures of tetracycline, tigecycline, and omadacycline [6].

activity over time has been examined by comparing the minimum inhibitory concentrations (MICs) from isolates collected during different surveillance periods. The MIC<sub>50/90</sub> values of omadacycline were similar for *S. pneumoniae* isolates collected in 2010 (0.06/0.12 µg/mL), 2014 (0.06/0.06 µg/mL), and 2016 (0.06/0.12 µg/mL) surveillance programs [12, 13]. Omadacycline has also remained highly active against clinical isolates of *S. aureus* across studies from 2010 (MIC<sub>90</sub> 0.25 µg/mL), 2014 (MIC<sub>90</sub> 0.12 µg/mL), and 2016 (MIC<sub>90</sub> 0.25 µg/mL) [13, 14]. Similarly, the MIC<sub>50/90</sub> values of omadacycline against isolates of *Legionella pneumophila* remained unchanged from 1995 (0.25/0.25 µg/mL) to 2014 (0.25/0.25 µg/mL) [15]. Omadacycline has been shown to be active against the category A biothreat pathogens *Bacillus anthracis* (MIC<sub>90</sub> 0.06 µg/mL) and *Yersinia pestis* (MIC<sub>90</sub> 1 µg/mL) [16]. Like other tetracyclines, omadacycline displays no notable activity against *Proteus* spp. (MIC<sub>90</sub> ≥32 µg/mL), *Providencia* spp. (MIC<sub>90</sub> >16 µg/mL), *Morganella* spp. (MIC<sub>90</sub> >16 µg/mL), or *Pseudomonas* spp. (MIC<sub>90</sub> >16 µg/mL) [13, 17]. In general, omadacycline has potent activity against atypical bacterial

pathogens and gram-positive aerobes, and a range of activities against gram-negative pathogens.

#### Gram-positive Aerobes

Omadacycline has potent in vitro activity against gram-positive aerobes, including antimicrobial-resistant pathogens such as MRSA (MIC<sub>90</sub> 0.25 µg/mL) and penicillin- or macrolide-resistant *S. pneumoniae* (MIC<sub>90</sub> 0.12 µg/mL; Table 1; breakpoints available from the US Food and Drug Administration [FDA]) [13, 18]. Omadacycline MIC values have been determined to be comparable for healthcare- and community-associated MRSA [14, 19]. Omadacycline is also active against VRE, with an MIC<sub>90</sub> of 0.25 µg/mL for vancomycin-resistant *Enterococcus faecalis* and of 0.12 µg/mL for vancomycin-resistant *Enterococcus faecium*. It also retains activity against tetracycline-resistant gram-positive bacteria, including *S. aureus* (MIC<sub>90</sub> 0.5 µg/mL), *E. faecalis* (MIC<sub>90</sub> 0.25 µg/mL), *E. faecium* (MIC<sub>90</sub> 0.12 µg/mL), *S. pneumoniae* (MIC<sub>90</sub> 0.12 µg/mL), and β-hemolytic streptococci (MIC<sub>90</sub> 0.25 µg/mL; Table 1) [13].

#### Gram-negative Aerobes

Omadacycline has in vitro activity against species of Enterobacteriaceae, including *Escherichia coli* (MIC<sub>90</sub> 2 µg/mL), *Klebsiella pneumoniae* (MIC<sub>90</sub> 8 µg/mL), *Klebsiella oxytoca* (MIC<sub>90</sub> 2 µg/mL), *Citrobacter* spp. (MIC<sub>90</sub> 4 µg/mL), and *Enterobacter cloacae* (MIC<sub>90</sub> 4 µg/mL; Table 2; breakpoints available from the FDA) [13, 18]. Omadacycline is also active against *Haemophilus influenzae* (MIC<sub>90</sub> 1 µg/mL; breakpoints available from the FDA) and *Moraxella catarrhalis* (MIC<sub>90</sub> 0.25 µg/mL; Table 2) [13, 18]. Omadacycline is not a substrate for extended-spectrum β-lactamases (ESBLs). It displays the same MIC<sub>90</sub> values against *E. coli* (2 µg/mL) and *K. pneumoniae* (8 µg/mL), whether ESBL negative or positive [20].

#### Atypical Bacteria and Anaerobes

Omadacycline displays potent in vitro activity against atypical bacteria, with an MIC<sub>90</sub> of 2 µg/mL for *Mycobacterium abscessus*; 0.5 µg/mL for *Mycobacterium fortuitum*; 0.25 µg/mL for *Mycobacterium chelonae*, *Mycoplasma pneumoniae*, *Chlamydia pneumoniae*, and *L. pneumophila*; and 0.06 µg/mL for *Mycoplasma hominis* (Table 3) [17, 21–23]. In addition, omadacycline has in vitro activity against anaerobic pathogens, including *Bacteroides fragilis* (MIC<sub>90</sub> 4 µg/mL), *Bacteroides thetaiotaomicron* (MIC<sub>90</sub> 4 µg/mL), *Bacteroides vulgatus* (MIC<sub>90</sub> 1 µg/mL), *Bacteroides ovatus* (MIC<sub>90</sub> 8 µg/mL), *C. difficile* (MIC<sub>90</sub> 0.5 µg/mL), *Clostridium perfringens* (MIC<sub>90</sub> 16 µg/mL), and anaerobic gram-positive cocci (MIC<sub>90</sub> 1 µg/mL; Table 4) [24].

#### FACTORS AFFECTING IN VITRO ACTIVITY

The in vitro activity of omadacycline against gram-positive and gram-negative clinical isolates was examined in the presence of

**Table 1. In Vitro Activity of Omadacycline and Comparators Against Select Gram-positive Aerobes**

Bacteria (No. of isolates)	Omadacycline				Tetracycline				Tigecycline			
	MIC <sub>50</sub> (µg/mL)	MIC <sub>90</sub> (µg/mL)	MIC Range (µg/mL)	% S by CLSI <sup>e</sup>	MIC <sub>50</sub> (µg/mL)	MIC <sub>90</sub> (µg/mL)	MIC Range (µg/mL)	% S by CLSI	MIC <sub>50</sub> (µg/mL)	MIC <sub>90</sub> (µg/mL)	MIC Range (µg/mL)	% S by CLSI
<i>S. aureus</i> (4215)	0.12	0.25	≤0.015 to 8	98.6	≤0.5	≤0.5	≤0.5 to 8	94.2	0.06	0.12	≤0.015 to 1	>99.9
<i>S. aureus</i> MSSA (2777)	0.12	0.25	≤0.015 to 1	99.9	≤0.5	≤0.5	≤0.5 to 8	96.3	0.06	0.12	≤0.015 to 0.25	100.0
<i>S. aureus</i> MRSA (1438)	0.12	0.25	0.03 to 8	96.1	≤0.5	4	≤0.5 to 8	90.3	0.06	0.12	≤0.015 to 1	99.9
<i>S. aureus</i> TR (221) <sup>a</sup>	0.12	0.5	0.03 to 2	95.5	–	–	–	–	0.12	0.25	0.03 to 1	99.5
<i>E. faecalis</i> (677)	0.12	0.25	≤0.015 to 1	97.2	>16	>16	≤0.12 to 16	21.4	0.06	0.12	≤0.015 to 0.12	100.0
<i>E. faecalis</i> VS (663)	0.12	0.25	≤0.015 to 1	97.1	>16	>16	≤0.12 to 16	21.7	0.06	0.12	≤0.015 to 0.12	100.0
<i>E. faecalis</i> VNS (14)	0.12	0.25	≤0.015 to 0.25	100.0	>16	>16	0.25 to 16	7.7	0.06	0.12	≤0.015 to 0.12	100.0
<i>E. faecalis</i> TR (524) <sup>a</sup>	0.12	0.25	≤0.015 to 1	96.4	–	–	–	–	0.06	0.12	≤0.015 to 0.12	100.0
<i>E. faecium</i> (390)	0.06	0.12	≤0.015 to 8	NA	16	>16	≤0.12 to 16	42.6	0.03	0.06	≤0.015 to 1	NA
<i>E. faecium</i> VS (234)	0.06	0.12	≤0.015 to 1	NA	0.5	>16	≤0.12 to 16	58.1	0.03	0.06	≤0.015 to 0.25	NA
<i>E. faecium</i> VNS (156)	0.06	0.12	≤0.015 to 8	NA	>16	>16	≤0.12 to 16	19.2	0.03	0.06	≤0.015 to 1	NA
<i>E. faecium</i> TR (217) <sup>a</sup>	0.12	0.12	≤0.015 to 8	NA	–	–	–	–	0.06	0.06	≤0.015 to 1	NA
<i>S. pneumoniae</i> (1314)	0.06	0.12	≤0.015 to 1	99.7	≤0.25	>8	≤0.25 to 8	79.5	0.03	0.06	0.015 to 0.25	99.4
<i>S. pneumoniae</i> PS (899)	0.06	0.06	≤0.015 to 0.5	99.9	≤0.25	0.5	≤0.25 to 8	92.2	0.03	0.06	0.015 to 0.12	99.4
<i>S. pneumoniae</i> PI (263)	0.06	0.12	≤0.015 to 1	98.9	0.5	>8	≤0.25 to 8	59.7	0.03	0.06	0.015 to 0.25	98.9
<i>S. pneumoniae</i> PR (152)	0.06	0.12	≤0.015 to 0.12	100.0	>8	>8	≤0.25 to 8	38.8	0.06	0.06	0.015 to 0.06	100.0
<i>S. pneumoniae</i> MR <sup>b</sup> (413)	0.06	0.12	≤0.015 to 1	99.5	>8	>8	≤0.25 to 8	44.1	0.06	0.06	0.015 to 0.25	99.5
<i>S. pneumoniae</i> TR (263) <sup>a</sup>	0.06	0.12	≤0.015 to 1	99.2	–	–	–	–	0.06	0.06	0.015 to 0.25	99.2
<i>S. anginosus</i> group (107)	0.06	0.12	≤0.015 to 0.12	100.0	0.5	>8	≤0.25 to 8	67.3	0.03	0.03	≤0.008 to 0.12	100.0
<i>S. anginosus</i> group TR (34) <sup>a</sup>	0.06	0.12	≤0.015 to 0.12	100.0	–	–	–	–	0.06	0.12	≤0.015 to 0.12	100.0
β-hemolytic streptococci <sup>c</sup> (866)	0.06	0.12	0.03 to 0.5	NA	0.5	>8	≤0.25 to 8	54.7	0.06	0.06	0.015 to 0.25	100.0
β-hemolytic streptococci TR (421) <sup>a</sup>	0.12	0.25	0.03 to 0.5	NA	–	–	–	–	0.06	0.06	0.015 to 0.25	100.0
β-hemolytic streptococci MR <sup>d</sup> (266)	0.12	0.25	0.03 to 0.5	NA	>8	>8	≤0.25 to 8	25.6	0.06	0.06	0.015 to 0.25	100.0

Adapted from Pfaler et al [13]; isolates were collected in the United States and Europe for the 2016 SENTRY Antimicrobial Surveillance Program.

Abbreviations: % S, percentage susceptible; CLSI, Clinical and Laboratory Standards Institute; *E. Enterococcus*; MIC, minimum inhibitory concentration; MR, macrolide-resistant; MRSA, methicillin-resistant *S. aureus*; MSSA, methicillin-susceptible *S. aureus*; NA, data not available; PI, penicillin-intermediate; PR, penicillin-resistant; PS, penicillin-susceptible; *S.*, *Staphylococcus*; TR, tetracycline-resistant; VNS, vancomycin-nonsusceptible; VS, vancomycin-susceptible.

<sup>a</sup>Isolates were defined as tetracycline-resistant by CLSI (M100-S27, 2017) breakpoint interpretive criteria; MIC<sub>50</sub> and MIC<sub>90</sub> data for tetracycline vs these isolates were not calculated.

<sup>b</sup>Includes erythromycin- and azithromycin-resistant *Streptococcus pneumoniae*.

<sup>c</sup>Includes *Streptococcus agalactiae*, *Streptococcus canis*, *Streptococcus dysgalactiae*, and *Streptococcus pyogenes*.

<sup>d</sup>Erythromycin-resistant β-hemolytic streptococci.

<sup>e</sup>Based on Food and Drug Administration breakpoints (see [18]); the ABSSSI breakpoint was used for *Staphylococcus aureus*.

**Table 2. In Vitro Activity of Omadacycline and Comparators Against Select Gram-negative Aerobes**

Bacteria (No. of Isolates)	Omadacycline				Tetracycline				Tigecycline			
	MIC <sub>50</sub> (µg/mL)	MIC <sub>90</sub> (µg/mL)	MIC Range (µg/mL)	% S by CLSI <sup>b</sup>	MIC <sub>50</sub> (µg/mL)	MIC <sub>90</sub> (µg/mL)	MIC Range (µg/mL)	% S by CLSI	MIC <sub>50</sub> (µg/mL)	MIC <sub>90</sub> (µg/mL)	MIC Range (µg/mL)	% S by CLSI
Enterobacteriaceae (8345)	1	8	0.12 to 32	NA	2	>16	≤0.25 to 16	64.2	0.25	0.25	≤0.06 to 8	97.8
Enterobacteriaceae CNS (1439)	2	8	0.12 to 32	NA	16	>16	0.5 to 16	44.5	0.25	0.25	≤0.06 to 8	98.2
Enterobacteriaceae TR (2737) <sup>a</sup>	2	16	0.12 to 32	NA	–	–	–	–	0.5	2	≤0.06 to 8	93.4
<i>E. coli</i> (3541)	0.5	2	0.12 to 32	NA	2	>16	≤0.25 to 16	63.7	0.12	0.25	≤0.06 to 4	99.9
<i>E. coli</i> /CS (3030)	0.5	2	0.12 to 32	NA	2	>16	≤0.25 to 16	69.1	0.12	0.25	≤0.06 to 2	100.0
<i>E. coli</i> CNS (511)	1	2	0.12 to 32	NA	>16	>16	0.5 to 16	31.7	0.12	0.25	≤0.06 to 4	99.6
<i>E. coli</i> TR (1272) <sup>a</sup>	1	4	0.12 to 32	NA	–	–	–	–	0.12	0.25	≤0.06 to 4	99.8
<i>K. pneumoniae</i> (1771)	2	8	0.25 to 32	89.7	2	>16	≤0.25 to 16	71.6	0.25	1	≤0.06 to 4	98.8
<i>K. pneumoniae</i> CS (1264)	1	4	0.25 to 32	94.7	1	>16	≤0.25 to 16	84.1	0.25	0.5	≤0.06 to 4	99.3
<i>K. pneumoniae</i> CNS (507)	2	8	0.25 to 32	77.1	16	>16	0.5 to 16	40.4	0.5	2	≤0.06 to 4	97.4
<i>K. pneumoniae</i> TR (430) <sup>a</sup>	4	16	0.5 to 32	67.7	–	–	–	–	0.5	2	≤0.06 to 4	94.9
<i>K. oxytoca</i> (423)	1	2	0.25 to 32	NA	1	8	≤0.25 to 16	89.8	0.25	0.5	≤0.06 to 2	100.0
<i>K. oxytoca</i> TR (30) <sup>a</sup>	2	16	0.25 to 32	NA	–	–	–	–	0.5	1	≤0.06 to 2	100.0
<i>E. cloacae</i> spp. complex (752)	2	4	0.25 to 32	93.6	2	16	0.5 to 16	85.6	0.25	0.5	0.12 to 4	99.2
<i>E. cloacae</i> spp. complex CS (542)	2	4	0.5 to 32	95.0	2	4	0.5 to 16	92.4	0.25	0.5	0.12 to 4	99.3
<i>E. cloacae</i> spp. complex CNS (210)	2	4	0.25 to 32	90.0	2	>16	1 to 16	68.1	0.5	1	0.12 to 4	99.0
<i>E. cloacae</i> spp. complex TR (87) <sup>a</sup>	4	16	1 to 32	69.0	–	–	–	–	0.5	2	0.12 to 4	95.4
Other <i>Enterobacter</i> spp. (250)	1	4	0.5 to 16	NA	1	4	0.5 to 16	91.2	0.25	0.5	0.12 to 4	99.6
Other <i>Enterobacter</i> spp. TR (17) <sup>a</sup>	16	16	2 to 16	NA	–	–	–	–	2	2	0.5 to 4	94.1
<i>Citrobacter</i> spp. (354)	1	4	0.25 to 16	NA	1	4	0.5 to 16	91.0	0.25	0.5	0.12 to 2	100.0
<i>Citrobacter</i> spp. TR (23) <sup>a</sup>	4	8	0.5 to 8	NA	–	–	–	–	0.5	1	0.12 to 2	100.0
<i>P. mirabilis</i> (463)	16	>32	2 to 32	NA	>16	>16	1 to 16	1.1	2	4	0.25 to 8	69.8
<i>P. mirabilis</i> TR (458) <sup>a</sup>	16	>32	2 to 32	NA	–	–	–	–	2	4	0.25 to 8	69.4
<i>Proteus</i> spp. IP (317)	8	32	0.5 to 32	NA	16	>16	0.5 to 16	38.2	1	2	0.12 to 8	97.2
<i>Proteus</i> spp. IP, TR (169) <sup>a</sup>	8	32	0.5 to 32	NA	–	–	–	–	1	2	0.12 to 8	95.3
<i>H. influenzae</i> (803)	1	1	0.12 to 16	99.4	0.5	1	≤0.06 to 8	99.8	0.12	0.25	0.06 to 1	96.1
<i>H. influenzae</i> BLP (201)	1	1	0.25 to 4	99.0	0.5	0.5	0.25 to 1	100.0	0.12	0.25	0.06 to 1	93.5
<i>H. influenzae</i> BLN (602)	1	1	0.12 to 16	99.5	0.5	1	≤0.06 to 8	99.7	0.12	0.25	0.06 to 1	97.0
<i>M. catarrhalis</i> (408)	0.25	0.25	0.06 to 0.5	NA	0.25	0.5	0.12 to 0.5	100.0	0.06	0.06	≤0.015 to 0.12	NA

Adapted from Pfaller et al [13]; isolates were collected in the United States and Europe for the 2016 SENTRY Antimicrobial Surveillance Program.

Abbreviations: % S, percentage susceptible; BLN, β-lactamase-negative; BLP, β-lactamase-positive; CLSI, Clinical and Laboratory Standards Institute; CNS, ceftazidime-nonsusceptible; CS, ceftazidime-susceptible; *E. cloacae*, *Enterobacter cloacae*; *E. coli*, *Escherichia coli*; *H. Haemophilus*; IP, indole-positive; *K. Klebsiella*; *M. Moraxella*; MIC, minimum inhibitory concentration; NA, data not available; *P. Proteus*; TR, tetracycline-resistant.

<sup>a</sup>Isolates were defined as tetracycline-resistant by CLSI (M100-S27, 2017) breakpoint interpretive criteria; MIC<sub>50</sub> and MIC<sub>90</sub> data for tetracycline vs these isolates were not calculated.

<sup>b</sup>Based on Food and Drug Administration breakpoints (see [18]).

**Table 3. In Vitro Activity of Omadacycline and Doxycycline Against Atypical Bacteria**

Bacteria (No. of Isolates)	Omadacycline			Doxycycline		
	MIC <sub>50</sub> (μg/mL)	MIC <sub>90</sub> (μg/mL)	MIC Range (μg/mL)	MIC <sub>50</sub> (μg/mL)	MIC <sub>90</sub> (μg/mL)	MIC Range (μg/mL)
<i>Mycoplasma hominis</i> (20)	0.03	0.06	0.016 to 0.12	0.06	2	0.016 to 2
<i>Mycoplasma pneumoniae</i> (20)	0.12	0.25	0.12 to 0.25	0.25	0.5	0.12 to 0.5
<i>Ureaplasma</i> spp. <sup>a</sup> (20)	1	2	0.25 to 2	0.25	4	0.06 to 4
<i>Legionella pneumophila</i> (100)	0.25	0.25	0.06 to 1	1	1	0.5 to 1
<i>Chlamydia pneumoniae</i> (15)	0.06	0.25	0.03 to 0.5	0.125	0.125	0.06 to 0.25
<i>Mycobacterium abscessus</i> (24)	1	2	0.06 to 8	>64	>64	0.25 to 64
<i>Mycobacterium chelonae</i> (22)	0.125	0.25	0.015 to 0.25	32	64	16 to 64
<i>Mycobacterium fortuitum</i> (20)	0.125	0.50	0.03 to 1	8	64	<0.06 to 64

Adapted from Kohlhoff et al [22], Villano et al [17], Waites et al [21], and Shoen et al [23].

Abbreviation: MIC, minimum inhibitory concentration.

<sup>a</sup>Includes *Ureaplasma parvum* (n = 10) and *Ureaplasma urealyticum* (n = 10).

either bovine lung surfactant or human or mouse serum [25]. As expected, MIC values of omadacycline against tetracycline-susceptible and tetracycline-resistant strains of *S. aureus*, *S. pneumoniae*, *H. influenzae*, and *E. coli* did not increase in the presence of surfactant or serum. In these assays, the addition of serum increased the MIC values of doxycycline, and the addition of surfactant increased the MICs of daptomycin. Studies examining the in vitro protein binding of omadacycline have shown that, over a concentration range of 10–10 000 ng/mL, omadacycline displays low binding to human (21%), monkey (21%), rat (26%), and mouse (15%) plasma proteins [26].

The pH is known to affect the activity of a range of antibiotics, because of its effect on the charge of the molecules [27]. Compared with standard pH 7.4 medium, omadacycline MIC values were unaffected by pH 8.0 medium, whereas MIC values were several-fold higher when tested at pH 5.0–6.0 [28]. In addition, in unpublished observations, omadacycline retained activity in pooled human urine (pH 6.6) and in pooled human urine adjusted to pH 7.1 (Paratek Pharmaceuticals, Inc.; data on file). In urine, omadacycline MIC values against *E. coli* and *Staphylococcus saprophyticus* were either unaffected or up to 2-fold higher than MICs observed in standard pH 7.3 broth

medium. Overall, these assays demonstrate that omadacycline retains activity in human urine.

Intracellular activity was evaluated for omadacycline against both *S. aureus* and *L. pneumophila*. In assays of intracellular activity, *S. aureus*-infected THP-1 human monocytes were treated with 1, 2, 8, or 16 times the MIC of either omadacycline or comparators (tigecycline, linezolid, ceftaroline, levofloxacin, moxifloxacin, or azithromycin) [29]. At 24 hours, omadacycline exhibited bactericidal activity against intracellular *S. aureus* (methicillin-susceptible and methicillin-resistant strains) with ≥99% growth reduction at 2–16 times the MIC, which was similar to that of levofloxacin and moxifloxacin and was more active than that of tigecycline (≥99% vs <99 to ≥90%), linezolid (≥99% vs <99 to ≥90%), ceftaroline (≥99% vs <90%), and azithromycin (≥99% vs <90%). Similar studies carried out with U937 human monocytes infected with erythromycin-susceptible or erythromycin-resistant strains of *L. pneumophila* serogroup 1 also demonstrated the robust intracellular activity of omadacycline [30].

These studies demonstrate that omadacycline possesses low binding to plasma proteins and that its activity is not significantly affected by surfactant, serum, high pH, or urine. The

**Table 4. In Vitro Activity of Omadacycline and Comparators Against Anaerobes**

Bacteria (No. of Isolates)	Omadacycline			Tigecycline		
	MIC <sub>50</sub> (μg/mL)	MIC <sub>90</sub> (μg/mL)	MIC Range (μg/mL)	MIC <sub>50</sub> (μg/mL)	MIC <sub>90</sub> (μg/mL)	MIC Range (μg/mL)
<i>Bacteroides fragilis</i> (21)	0.5	4	0.25 to 16	0.5	2	0.5 to 8
<i>Bacteroides thetaiotaomicron</i> (21)	1	4	0.12 to 16	1	8	0.25 to 16
<i>Bacteroides vulgatus</i> (21)	0.12	1	0.06 to 2	0.25	1	0.12 to 2
<i>Bacteroides ovatus</i> (15)	0.5	8	0.06 to 16	0.5	8	0.03 to 16
<i>Clostridioides difficile</i> (21)	0.25	0.5	0.25 to 8	0.25	0.25	0.25 to 4
<i>Clostridium perfringens</i> (22)	4	16	0.12 to 16	8	>16	0.25 to 16
<i>Peptostreptococcus</i> spp. <sup>a</sup> (22)	0.12	1	0.06 to 2	0.12	2	0.06 to 4

Adapted from Villano et al [17] and Stapert et al [24].

Abbreviation: MIC, minimum inhibitory concentration.

<sup>a</sup>Includes *Peptostreptococcus micros* (n = 11) and *Peptostreptococcus anaerobius* (n = 11).

human monocyte models demonstrate the intracellular penetration of active omadacycline, which has clinical implications for the treatment of pneumonia.

## IN VITRO PHARMACODYNAMIC PROPERTIES

The in vitro pharmacodynamic properties of omadacycline have been studied in a variety of ways. Omadacycline displays bactericidal or bacteriostatic activity that is organism-dependent. In a study examining 85 bacterial isolates, minimum bactericidal concentrations indicated that omadacycline has bactericidal activity ( $\geq 3$  log reduction of initial inoculum) against *S. pneumoniae*, *H. influenzae*, and *M. catarrhalis*, and that it displays bacteriostatic activity against enterococci, *S. aureus*, and most isolates of *E. coli* [31].

In order to determine the residual activity of omadacycline after antibiotic removal, the postantibiotic effect (PAE) was evaluated. The growth inhibition of omadacycline following drug removal has been examined for clinical isolates of *S. aureus* (including 1 MRSA isolate), *S. pneumoniae* (including 1 penicillin-resistant isolate), enterococci (including 1 vancomycin-resistant isolate), and *E. coli* [32]. For these gram-positive and gram-negative aerobes, omadacycline had a PAE ranging from 1.4 to 3.3 hours (1 hour initial exposure at 5 times the MIC), which was similar to that of tigecycline; with the exception of enterococci, for which tigecycline displayed longer PAEs [32]. The activity of omadacycline against *E. coli* biofilms was determined with the minimum biofilm eradication concentration assay (Innovotech, Edmonton, Alberta, Canada) [33]. Omadacycline displayed dose-dependent activity against established biofilms, with a reduction of  $\sim 2$  log<sub>10</sub> units in biofilm-associated bacteria. At concentrations near the MIC (1.13  $\mu\text{g}/\text{mL}$ ), omadacycline significantly reduced the total *E. coli* bioburden. In addition, *E. coli* biofilms were not propagated in the presence of sub-MIC concentrations of omadacycline. This finding may have clinical implications, as biofilm colonies are known to be resistant to sub-MIC doses of most antibiotics [34].

The effect of omadacycline on the gut microflora and the potential for omadacycline to induce *C. difficile* infections (CDI) have been investigated in an in vitro human gut model [35]. In this model, the bacterial compositions of the proximal, medial, and distal colon are simulated with a 3-stage continuous system of temperature- and pH-controlled vessels [36]. Antimicrobials that are considered high risk for CDI have been shown to induce simulated CDI in this in vitro human gut model, whereas antimicrobials that are considered low risk for CDI have not induced simulated CDI in this model [36–39]. The 3-vessel system was equilibrated following inoculation with pooled feces from healthy volunteers, and then infected with *C. difficile* spores, followed by omadacycline treatment [35]. Although omadacycline disrupted the gut microflora (declines in Bifidobacteria, *B. fragilis* group spp., *Lactobacillus* spp.,

*Enterococcus* spp., *Clostridium* spp., and Enterobacteriaceae populations), there was no evidence of simulated CDI (ie, *C. difficile* germination, vegetative cell proliferation, or toxin production). In contrast, moxifloxacin disrupted the gut microflora and induced simulated CDI in this study [35], which supports the clinical finding that fluoroquinolones have one of the highest incidences of CDI [40].

## IN VIVO PHARMACODYNAMIC PROPERTIES

The in vivo pharmacodynamics of omadacycline have been examined in 2 animal models: a neutropenic mouse model of pneumonia and a neutropenic mouse thigh infection model [41, 42] (Table 5). The primary pharmacokinetic/pharmacodynamic target predictive of efficacy for omadacycline was noted to be the ratio of the area under the plasma concentration–time curve over 24 hours to the MIC (AUC/MIC). Omadacycline AUC/MIC correlated with microbiological efficacy ( $r^2 = 0.74$ ; mean plasma 24-hour static dose AUC/MIC = 16–20) in mice infected in the lungs with *S. pneumoniae*, including those with strains with varying susceptibility to tetracyclines,  $\beta$ -lactams, and macrolides [41]. Based on omadacycline concentration measurements in plasma and epithelial lining fluid, omadacycline penetrated well into epithelial lining fluid [41]. In the mouse thigh infection model (which included 10 isolates of *S. aureus*, including MRSA), the efficacy of omadacycline was defined by the AUC/MIC ( $r^2 = 0.92$ ; mean plasma 24-hour static dose AUC/MIC = 24) [42]. These findings are in agreement with those for tetracyclines, as the AUC/MIC ratio is the pharmacodynamic parameter that best correlates with treatment efficacy for this class [43].

## IN VIVO EFFICACY

The in vivo efficacy of omadacycline has been demonstrated in a number of animal models (Table 5). Omadacycline, administered as a single intravenous dose, displayed potent efficacy in a mouse model of intraperitoneal infection with *E. coli* or tetracycline-susceptible and tetracycline-resistant strains of *S. aureus* and *S. pneumoniae* [8] (Table 5). Overall in this model, the efficacy of omadacycline was similar to, or greater than, that of comparators. In a mouse intra-abdominal infection model of postoperative polymicrobial peritonitis, 2 intravenous doses of omadacycline showed increased 10-day survival over comparators [44] (Table 5). In a urinary tract infection model, mice were infected with *E. coli* directly into the bladder and then given increasing single doses of omadacycline on day 4 postinfection. Omadacycline performed as well as minocycline (50% effective dose, 4.3 vs 4.5 mg/kg) in this model [45] (Table 5). Against the biothreat pathogens—*B. anthracis* and *Y. pestis*—omadacycline displayed in vivo efficacy in murine whole-body aerosol infection models [16] (Table 5).

**Table 5. Efficacy of Omadacycline in Animal Models**

Study	Animal Model	Strain	Antimicrobial	Result
Lepak et al [41]	Neutropenic mouse pneumonia model (female ICR/Swiss mice, 3 mice/group)	<i>Streptococcus pneumoniae</i> <sup>a</sup> 1293 ATCC 10813	SC administration <sup>b</sup> Omadacycline	Plasma AUC/MIC $E_{max}$ 4.85 $ED_{50}$ 15.60 $r^2$ 0.74
Lepak et al [42]	Mouse neutropenic thigh infection model (4 thigh infections/group)	140 ATCC 49619	Omadacycline	Epithelial lining fluid AUC/MIC $E_{max}$ 4.91 $ED_{50}$ 15.11 $r^2$ 0.75
		<i>Staphylococcus aureus</i> MSSA	SC administration <sup>c</sup> Omadacycline	Thigh bacterial burden Reduction of 4–5 log <sub>10</sub> CFU/thigh compared with untreated controls
		<i>S. aureus</i> MRSA	Omadacycline	Plasma AUC/MIC $E_{max}$ 4.62 $ED_{50}$ 21.7 $r^2$ 0.92
Macone et al [8]	Mouse IP infection model (male CD-1 mice, 5 mice/group)	<i>S. pneumoniae</i> PBS1339 <i>S. pneumoniae</i> 700905 <i>S. aureus</i> 29213 <i>S. aureus</i> USA300 <i>S. aureus</i> MRSA5 <i>Escherichia coli</i> PBS1478	IV administration <sup>d</sup> Omadacycline Tigecycline Omadacycline Tigecycline Omadacycline Tigecycline Omadacycline Tigecycline Omadacycline Tigecycline	MIC (µg/mL) 0.125 0.125 ≤0.06 0.125 0.25 0.125 0.25 0.125 0.25 ≤0.06 ≤0.06 ED <sub>50</sub> (mg/kg [95% CI]) 3.34 ± 1.56 4.13 (2.46 to 5.79) 0.45 (0.32 to 0.58) 1.72 (0.6 to 2.82) 1.74 (0.91 to 2.58) 0.73 (0.69 to 0.76) 0.90 (0.33 to 1.46) 0.58 (0.40 to 0.75) 0.30 (0.295 to 0.305) 1.74 (0.91 to 2.57) 2.02 (1.09 to 2.96) 1.75 (1.12 to 2.38)
Endermann et al [44]	Mouse postoperative polymicrobial peritonitis model (female CFW-1 mice, 10 mice/group)	<i>Enterococcus faecalis</i> TR <i>Enterococcus faecium</i> VR	IV administration <sup>f</sup> Omadacycline Imipenem Linezolid	10-day survival 80% 70% 30%
McKenney et al [45]	Mouse urinary tract infection model (male CD-1 mice)	<i>E. coli</i> C189P4	IV administration <sup>g</sup>	MIC (µg/mL) Kidney bacterial burden (ED <sub>50</sub> mg/kg) 0.5 4.3 0.5 4.5 ≤0.06 <1.0
Steenbergen et al [16]	Mouse whole-body aerosol infection model, delayed treatment exposure (female BALB/c mice, 9 or 10 mice/group)	<i>Bacillus anthracis</i> Ames	IP administration <sup>h</sup> Omadacycline Doxycycline Ciprofloxacin Vehicle	40-day survival 60% 70% 80% 0%

**Table 5. Continued**

Study	Animal Model	Strain	Antimicrobial	Result
Kim et al [46]	Hamster <i>C. difficile</i> infection model (male LGV Golden Syrian hamsters, 10 hamsters/group)	<i>Yersinia pestis</i> CO92	IP administration <sup>1</sup> Omadacycline Doxycycline Ciprofloxacin Vehicle	40-day survival 90% (40 mg/kg dose group) 90% (40 mg/kg dose group) 100% 0%
		<i>Clostridioides difficile</i> ATCC 43596	Oral administration <sup>1</sup> Omadacycline Vancomycin	Median survival (days) 12 ( $P = .0004$ ) 2 ( $P = .0293$ )

Abbreviations: AUC, area under the plasma concentration–time curve; CFU, colony-forming units; CI, confidence interval; ED<sub>50</sub>, 50% effective dose; E<sub>max</sub>, maximum effect; IP, intraperitoneal; IV, intravenous; MIC, minimum inhibitory concentration; MRSA, methicillin-resistant *Staphylococcus aureus*; MSSA, methicillin-susceptible *Staphylococcus aureus*; SC, subcutaneous; TR, tetracycline-resistant; VR, vancomycin-resistant.

<sup>1</sup>There were 4 strains examined, with varying susceptibility to tetracyclines, β-lactams, and macrolides.

<sup>2</sup>There were 4-fold increasing doses examined, from 0.1 to 25.6 mg/kg.

<sup>3</sup>Doses increased 4-fold every 12 hours, from 0.25 to 64 mg/kg.

<sup>4</sup>At least 4 dose levels per experiment, with doses typically ranging from 0.11 to 18 mg/kg (dose minimum–maximum, 0.08–54 mg/kg); only tigecycline comparator is shown.

<sup>5</sup>Data are represented as means ± standard deviations from 7 independent experiments.

<sup>6</sup>Two doses of 10 mg/kg administered at 4 hours and 18 hours postsurgery.

<sup>7</sup>Increasing single doses on day 4 postinfection.

<sup>8</sup>Omadacycline: 15 mg/kg; doxycycline: 15 mg/kg; ciprofloxacin: 30 mg/kg; vehicle: 0.2 mL saline. All treatments began 48 ± 1 hours postinfection, and were given twice daily for 14 days.

<sup>9</sup>Omadacycline: 5, 10, 20, or 40 mg/kg; doxycycline: 5, 10, 20, or 40 mg/kg; ciprofloxacin: 15 mg/kg; vehicle: 0.2 mL saline. All treatments began 24 ± 1 hours postinfection, and were given twice daily for 7 days.

<sup>10</sup>Given for 5 days at 50 mg/kg/day.

The in vivo efficacy of omadacycline has also been examined in a hamster model of CDI [46] (Table 5). In this model, CDI was induced in hamsters by the subcutaneous administration of clindamycin 24 hours before infection with *C. difficile* by oral gavage. At 24 hours postinfection, hamsters were treated with oral omadacycline for 5 days. Omadacycline demonstrated efficacy in this model, with overall median survival of 12 days in omadacycline-treated hamsters compared with 2 days in vancomycin-treated hamsters.

In these in vivo models, omadacycline demonstrated efficacy greater than, or similar to, comparator antimicrobials.

## CONCLUSIONS

Unlike older tetracyclines, omadacycline is active against bacterial isolates that express tetracycline-specific efflux pumps and/or ribosomal protection resistance mechanisms. The in vitro antimicrobial activity of omadacycline covers a wide range of gram-positive and many gram-negative pathogens, including MRSA, penicillin- and macrolide-resistant *S. pneumoniae*, β-hemolytic streptococci, VRE, and Enterobacteriaceae, such as *E. coli* and *Klebsiella* spp. In addition, omadacycline displays in vitro antimicrobial activity against atypical and anaerobic organisms, including *Mycoplasma* spp., *L. pneumophila*, *Ureaplasma* spp., and *C. difficile*.

Omadacycline remains active at pH 8.0 and in the presence of serum, lung surfactant, or urine. In addition, omadacycline is active against bacterial biofilms and does not propagate biofilm formation. It has low binding to plasma proteins, displays a PAE of 1.4–3.3 hours (at 5 times the MIC, 1-hour exposure), and demonstrates intracellular activity against *S. aureus* and *L. pneumophila*.

In vivo pharmacodynamic studies have shown that, similar to other tetracyclines, omadacycline AUC/MIC has the strongest correlation with bacteriological outcome. The efficacy of omadacycline has been validated in a human gut model of *C. difficile* infection (in vitro) and in animal models (in vivo), including models of pneumonia, thigh infection, systemic infection, intra-abdominal infection, urinary tract infection, and CDI. Taken together, these studies demonstrate the activity of omadacycline against bacterial pathogens commonly associated with serious, community-acquired bacterial infections, including infections of the skin, lungs, and urinary tract.

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