Activity of JNJ-Q2 against *Staphylococcus aureus* Isolated from Patients with Acute Bacterial Skin and Skin-Structure Infection Obtained During a Phase II Clinical Trial

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Amended Abstract

Objective: To determine the activity of JNJ-Q2 against *S. aureus* isolated from patients with clinically diagnosed acute bacterial skin and skin-structure infection (ABSSSI) in the United States (USA) during a Phase II clinical trial and to determine the mechanisms of fluoroquinolone (FQ) resistance (R) in FQ-R strains. JNJ-Q2 is a broad-spectrum bactericidal 4-fluoroquinolone with potent activity against Gram-positive and -negative pathogens.

Materials and Methods

<u>Bacterial Strain Collection</u>. During the Phase II clinical trial, a total of 111 baseline *S. aureus* isolates were obtained from 111 patients, diagnosed with ABSSSI by strict criteria, in 15 medical centers throughout the United States (USA). Species identifications were performed by the submitting laboratories with confirmation performed by the central reference laboratory (JMI Laboratories, North Liberty, Iowa, USA).

Susceptibility Test Methods. All isolates were

Table 1. MIC frequency and cumulative percent inhibited distributions of JNJ-Q2 tested against 111 *S. aureus*^a.

Organism	No. (cum. %) of isolates inhibited at JNJ-Q2 MIC (mg/L)							
(no. tested)	≤0.004	0.008	0.015	0.03	0.06	0.12	0.25	
S. aureus (111)	27	25	7	1	0	49	2	
S. aureus (111)	(24.3)	(46.9)	(53.2)	(54.1)	(54.1)	(98.2)	(100.0)	
MSSA (42)	12	2	6	1	0	10	1	
1000A (42)	(28.6)	(57.2)	(71.4)	(73.8)	(83.0)	(97.6)	(100.0)	
MRSA (69)	15	13	1	0	0	39	1	
	(21.7)	(40.6)	(42.0)	(42.0)	(42.0)	(98.6)	(100.0)	
a. Isolates from a Pha	ase II clinical tri	al.						

Organism		CLSIb			
(no. tested)/	MIC ₅₀	MIC ₉₀	Range	%S / %R	
Antimicrobial agent ^a S. aureus – all isolates			range	7007 7010	
JNJ-Q2	0.015	0.12	≤0.004 – 0.25	_c / _	
Moxifloxacin	0.25	2	≤0.004 - 0.25	54.1 / 31.5	
Levofloxacin	2	4	<u>≤</u> 0.00 – 8 0.12 – >8	47.8 / 46.9	
Oxacillin	>2	4 >2	≤0.25 – >2	37.8 / 62.2	
Erythromycin	>8	>2 >8	<u> </u>	35.1 / 64.9	
Clindamycin	<i>≥</i> 0 ≤0.12	≥0 ≤0.12	≤0.12 – >8	97.3 / 2.7	
Linezolid	<u>−</u> 0.12 1	<u> </u>	≤0.12 - >0 1 - 2	100.0 / 0.0	
TMP/SMX ^d	ı ≤0.5	∠ ≤0.5	≤0.5 – >4	94.6 / 5.4	
	≤0.3 ≤0.25	≤0.5 0.5	≤0.3 – <i>></i> 4 ≤0.25 – 0.5	94.075.4 100.07-	
Daptomycin Vancomycin	≤0.25 1	0.5	≤0.25 – 0.5 0.5 – 2	100.0 / 0.0	
•	I		0.5 - 2	100.070.0	
MRSA (69) JNJ-Q2	0.12	0.12	≤0.004 – 0.25	- / -	
Moxifloxacin	0.12	2	≤0.004 – 0.25 ≤0.06 – 8	42.0 / 40.6	
	1	2 8	≤0.00 – 8 0.12 – >8	40.6 / 59.4	
Levofloxacin	4	-			
Erythromycin	>8	>8	0.25 ->8	15.9/84.1	
Clindamycin	≤0.12	≤0.12 2	≤0.12 – >8	95.7 / 4.3	
		2	1 - 2	100.0 / 0.0	
TMP/SMX ^d	≤0.5 <0.25	≤0.5 0.5	≤0.5 – 1	100.0 / 0.0	
Daptomycin	≤0.25	0.5	≤0.25 – 0.5	100.0 / -	
Vancomycin	I	I	0.5 – 2	100.0 / 0.0	
MSSA (42)	0.000	0.40		1	
JNJ-Q2 Mayiflayaasin	0.008	0.12	≤0.004 – 0.25	-/-	
Moxifloxacin	≤0.06 0.25	2	≤0.06 – 8 0.12 → 8	73.8 / 16.7	
Levofloxacin	0.25	4	0.12 ->8	59.5 / 26.2	
Erythromycin	0.5	>8	0.25 ->8	66.7 / 33.3	
Clindamycin	≤0.12 1	≤0.12 2	≤0.12 – 0.25	100.0/0.0	
		2	1 - 2	100.0 / 0.0	
TMP/SMX ^d	≤0.5 <0.25	>4	≤0.5 – >4	85.7 / 14.3	
Daptomycin Vancomycin	≤0.25	≤0.25	≤0.25 – 0.5 0.5 – 2	100.0 / - 100.0 / 0.0	

Methods: During the Phase II clinical trial, a total of 111 baseline *S. aureus* isolates were obtained from 111 patients, diagnosed with ABSSSI by strict criteria (including 37.8% methicillin-susceptible [MSSA] and 62.2% methicillin-resistant [MRSA]). Susceptibility testing was performed by the CLSI broth microdilution. Type II topoisomerase quinolone-resistant determinant regions (QRDR) were amplified by PCR and sequenced for FQ-R strains.

Results: JNJ-Q2 demonstrated good activity against all S. aureus and was very active against both MSSA (MIC_{50/90}, 0.008/0.12 mg/L) and MRSA (MIC_{50/90}, 0.12/0.12 mg/L). 51 strains (45.9%) had moxifloxacin MIC values of $\geq 1 \text{ mg/L}$ (non-susceptible) and all of these strains carried at least one *par*C mutation (S80Y). Additionally, 43 of 51 had gyrA S84L mutations and 3 strains also had *par*C E84G (1 strain), E84I (1), or *par*E T461I (1). Furthermore, 49 of 51 strains had a JNJ-Q2 MIC of 0.12 mg/L (range for 51, 0.12-0.25 mg/L). All isolates were susceptible to linezolid (LZD) and vancomycin (VAN). JNJ-Q2 was the most active agent tested with a MIC₉₀ 16-, 64- 16-, and eight-fold lower than MOX, levofloxacin (LEV), LZD and VAN, respectively.

tested for susceptibility by reference broth microdilution methods using the Clinical and Laboratory Standards Institute recommendations (CLSI; M07-A9, 2012) in validated panels manufactured by ThermoFisher Scientific Inc, formerly TREK Diagnostics Systems (Cleveland, Ohio, USA). Categorical interpretation of comparator MIC values was performed according to CLSI (M100-S22, 2012) and EUCAST (2012) criteria. The quality assurance of the minimum inhibitory concentration (MIC) values was performed by concurrent testing of CLSIrecommended (M100-S22, 2012) control strains, including *S. aureus* ATCC 29213 and Enterococcus faecalis ATCC 29212. The inoculum density was monitored by colony counts to assure an adequate number of cells for each testing event. All QC results were within established ranges.

<u>Molecular Methods</u>. QRDR mutations were detected by PCR amplification and sequencing

Conclusions

= No breakpoint has been established

d. Trimethoprim-sulfamethoxazole

 In summary, JNJ-Q2 was found to be very active against *S. aureus* isolated from patients with a definitive clinical diagnosis of ABSSSI during a 2010 Phase II clinical trial in the USA, with all 111 baseline strains inhibited at JNJ-Q2 MIC values of ≤0.25 mg/L. Additionally, JNJ-Q2 was the most active compound with activity being manyfold higher than the comparator agents tested.

Conclusions: JNJ-Q2 demonstrated very potent activity against contemporary *S. aureus* isolated from patients in the USA with clinically diagnosed and microbiologically confirmed ABSSSIs. JNJ-Q2 exhibited greater activity compared to LEV and MOX, including strains R to currently utilized FQs. These encouraging results support the further clinical development of JNJ-Q2 for ABSSSIs.

Introduction

Quinolone resistance in Staphylococcus

of *gyr*A, *gyr*B, *par*C and *par*E as previously described (Horii, et al., 2003). Sequences were aligned against *S. aureus* ATCC 25923 using MegAlign (DNAStar, Lasergene, Madison, Wisconsin, USA).

Results

- Of the 111 S. aureus isolated, 37.8% were methicillin-susceptible (MSSA) and 62.2% methicillin-resistant (MRSA). JNJ-Q2 demonstrated good activity against all isolates and was very active against both MSSA (MIC_{50/90}, 0.008/0.12 mg/L) and MRSA (MIC_{50/90}, 0.12/0.12 mg/L) with all isolates inhibited at JNJ-Q2 MIC values of ≤0.25 mg/L (Table 1).
- Levofloxacin and moxifloxicin resistance rates were high at 46.9 and 31.5%, respectively (Table 2). Levofloxacin/ moxifloxicin resistance rates were much higher in MRSA (59.4/40.6%) compared to
- These favorable results, along with the previously demonstrated low propensity for pathogens to induce rapid mutational resistance to JNJ-Q2, support the further clinical development of JNJ-Q2 to treat ABSSSI, including MRSA.

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aureus results from either mutations in the quinolone-resistance determinant regions (QRDRs) of the target enzymes, most commonly DNA gyrase and topoisomerase IV, or by drug efflux and/or decreased uptake. JNJ-Q2, a novel fluorinated 4-quinolone, has been shown to have balanced potency against both DNA gyrase and topoisomerase IV, excellent *in vitro* activity against both methicillin-susceptible *S. aureus* (MSSA) and methicillin-resistant *S. aureus* (MRSA), and is in clinical development for the treatment of acute bacterial skin and skin-structure infection (ABSSSI).

The aims of this study were to determine comparative *in vitro* activity for JNJ-Q2 tested against *S. aureus* obtained from a recent (2010) randomized, controlled, double-blind, double-dummy, multicenter Phase II study for JNJ-Q2 compared with linezolid for the treatment of ABSSSI (trial #NCT01128530). MSSA (26.2/16.7%), respectively. JNJ-Q2 was the most active agent tested with a MIC_{90} 16-, 32-, 16-, and eight-fold lower than moxifloxacin, levofloxacin, linezolid and vancomycin, respectively.

- 51 strains (45.9%) had moxifloxacin MIC values of ≥1 mg/L (non-susceptible) and all of these strains carried at least one *par*C mutation (S80Y). Additionally, 43 of 51 had *gyr*A S84L mutations and 3 strains also had *par*C E84G (1 strain), E84I (1), or *par*E T461I (1). Furthermore, 49 of 51 strains had a JNJ-Q2 MIC of 0.12 mg/L (range for 51, 0.12-0.25 mg/L).
- Overall erythromycin resistance was also elevated at 64.9%. Most isolates were susceptible to clindamycin (97.3%) and trimethoprim-sulfamethoxazole (94.6%). All isolates were susceptible to linezolid and vancomycin.