Identification and Susceptibility of Pathogens Isolated from Patients with

Complicated Skin and Skin Structure Infections (cSSSI): Results of a PTK796 (PTK) Phase 2 Clinical Trial

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ABSTRACT

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Background Paratek conducted a randomized, investigator-blinded Phase 2 clinical trial in adults with cSSSI comparing PTK (100mg IV/200mg oral, QD) with linezolid (LZD) (600mg IV/600 mg oral, Q12h). LZD was supplemented with aztreonam when gram negative infection was suspected. A total of 234 patients were enrolled.

Methods Patient enrollment isolates were identified and tested at a central laboratory. MICs were determined by broth microdilution using dried panels. Analysis was based on the Clinically Evaluable (CE) population with a pathogen isolated at enrollment (ME).

Results The predominant pathogen isolated was S. aureus (SA). 67 patients in the PTK arm were clinical successes with 80 isolates of SA (56.3% MRSA) and 59 isolates from 45 clinical successes (67.8% MRSA) in the LZD arm. MIC_{or}s for MRSA and MSSA from each treatment arm were 0.25 µg/ml (PTK) and 2.0 μg/ml (LZD). Other pathogens in the PTK clinical success group included β-hemolytic strep (4 pts, mixed cultures), Gram negative bacteria (5 pts, 4 mixed), E. faecalis (2pts, one mixed), S. hemolyticus (1pt, mixed), and S. constellatus (1pt, only pathogen). In the LZD clinical success group other pathogens included Gram negatives (4 pts., 3 mixed), E. faecium (1pt, mixed), and S. constellatus (2 pts, only pathogen). There were only two clinical failures in the PTK arm, both MSSA (PTK MIC = 0.25 and 0.5 ug/ml, mixed cultures) and three in the LZD arm. 2 MRSA (LZD MIC = 2ug/ml. 1 mixed) and one MSSA (LZD MIC = 2ug/ml. only

Conclusions PTK exhibited excellent efficacy in the Phase 2 trial compared to LZD (CE Success 98% vs 93.2% respectively and ME Success 97.4% vs 93.7%, respectively). S. aureus was the predominant pathogen at enrollment (93.3% from PTK and 87.3% from LZD) the majority of which were MRSA. Susceptibility to PTK and LZD did not differ between MRSA and MSSA and did not differ between treatment arms.

INTRODUCTION

PTK796 is a novel aminomethylcycline currently in worldwide development by Paratek Pharmaceuticals and Novartis Pharmaceuticals. PTK796 is a member of the tetracycline family and exhibits excellent in vitro activity and efficacy in animal infection models. PTK796 is active against the most common pathogens associated with skin and skin structure infections, most notably community acquired MRSA and betahemolytic streptococci. PTK796 overcomes tetracycline resistance common to these pathogens, either tetracycline efflux or ribosomal protection. Paratek Pharmaceuticals conducted a Phase 2 clinical trial in complicated skin and skin structure infection from July 2007 through January 2008. The study was a randomized, investigator blinded study comparing intravenous and oral PTK796 to linezolid in the treatment of complicated skin and skin structure infections (1).

STUDY DESIGN

The Phase 2 trial was an investigator blinded evaluation in which patients with qualifying infections and clinical status were randomized to receive either 100 mg PTK796 intravenously OD or 600 mg linezolid intravenously BID. In cases suspected of being caused by Gram negative pathogens, patients received, in blinded fashion), either placebo (saline) infusions (PTK796 treatment group) or aztreonam (linezolid treatment group). Following the intravenous course and at the decision of the principal investigator, patients were discharged with either 200 mg oral PTK796 OD or 600mg linezolid. BID.

Patient demographics were well balanced between the two treatment groups. As shown in Table 1 the types of infection were similar between groups although infections tended to be larger and more severe in the PTK796 treatment group(1). Clinical outcome are shown in Table 2.

Table 1. Characteristics of Infactions (ITT)

Type of Infection	N	Median (cm) Maximum Dimension (range)	N Median (cm) Maximum Dimension (rang	
		PTK796	Linezolid	
Major Abscess	73	10 (6-16)	72	7.8 (4.1-13)
Injury	21	10.5 (4-31)	17	7 (3-19.5)
Cellulitis	8	20 (12-31)	10	18 (3.3-33)
Lower Extremity Ulcer	9	3.5 (1.2-10)	9	2.5 (1 - 28)

Table 2. Derivation of Patient Population

Population ¹	Treatment	Total #	Patients by Clinical Outcome ²		
	Group	Patients	Success	Failure	
ITT	PTK796	111	98	2	
	Linezolid	108	82	6	
mITT	PTK796	84	75	2	
	Linezolid	78	59	4	
Clinically Evaluable	PTK796	100	98	2	
(CE)	Linezolid	88	82	6	
Microbiologically	PTK796	77	75	2	
Evaluable (ME)	Linezolid	63	59	4	
ME w/	PTK796	71	69	2	
Susceptibility (MES)	Linezolid	49	46	3	

ITT= All patients enrolled and treated; mITT=All patients with at least one pathogen isolated at baseline; CE=all ITT patients completing all clinical evaluations; ME = all CE pts with at least one pathogen at baseline; ME w/Susceptibility= All ME with central

2Patients not completing evaluations for administrative reasons have been excluded

MICROBIOLOGY METHODS

Qualified samples of infected tissue were processed by local clinical microbiology laboratories. Isolated organisms were sent to a central laboratory for definitive identification and susceptibility testing. Susceptibility to PTK796 was determined using dried MIC panels (TREK Diagnostics, Cleveland, OH) previously shown to perform comparably to standard reference methods. Linezolid, oxacillin, and aztreonam susceptibilities were determined using commercially available methods. The Primary population for this report is the clinically evaluable population for which there was at least one pathogen at baseline for which an MIC was determined. In some cases, there were multiple isolates with slightly different MICs that were included in the analyses.

RESULTS

As shown in Table 3. Staphylococcus aureus was the most frequently isolated pathogen (67/71 patients in the PTK796 treatment group and 45/49 in the linezolid treatment group. It was most often the sole pathogen isolated, but did occur in mixed infections.

Table 3, Types of Pathogens at Baseline

Organism	Number of Isolates				
	PTK796 Linezolid (71 patients, 67 with (49 patients, 45 wi S. aureus) S.aureus)				
MRSA	45	40			
MSSA	35	19			
β-hemolytic Streptococci	5	2			
Enterococcus sp	4	1			
Other Gram + Cocci	2	1			
Gram Negative Bacilli	14	5			

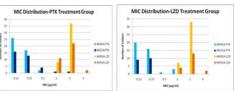
The susceptibility of the isolates to study antibiotics is shown in Table 4. PTK796 exhibited potent activity against S. aureus regardless of whether the organism was MRSA or MSSA.

Table 4. Susceptibility of S. aureus to study antibiotics

Organism	Treatment	#	Antibiotics	1)	
	Group	Isolates		MIC Range	MIC ₅₀	MIC ₉₀
S. aureus	PTK796	80	PTK796	0.12-2	0.12	0.25
			Linezolid	1-4	2	2
	Linezolid	59	PTK796	0.12-1	0.25	0.25
			Linezolid	1-4	2	2
MSSA	PTK796	35	PTK796	0.12-2	0.25	0.5
			Linezolid	1-4	2	2
	Linezolid	19	PTK796	0.12-0.25	0.12	0.25
			Linezolid	1-4	2	4
MRSA	PTK796	45	PTK796	0.12-0.5	0.12	0.25
			Linezolid	1-2	2	2
	Linezolid	40	PTK796	0.12-1	0.12	0.25
			Linezolid	1-2	2	2

The distribution of PTK and linezolid MICs for S. aureus are shown in Figure 1a and 1b.

Figure 1a, MIC Distribution for S. aureus Figure 1a, MIC Distribution for S. aureus (PTK796 Treatment Group)



(Linezolid Treatment Group)

There were 2 clinical failures in the PTK796 treatment group and 3 clinical failures in the linezolid treatment group, all with S. aureus. In each case there was no indication that susceptibility was a factor in the clinical outcome since MICs were consistent with the entire population (Table 5).

Table 5 Identification and Susceptibility of Courses from Clinical Follows at

Treatment Group	Patient # Organism at Baseline		MIC (μg/ml)	
			PTK796	Linezolid
PTK796	#1	MSSA	0.5	2
	#2	MRSA	0.25	2
Linezolid	#1	MRSA	0.5	2
	#2	MSSA	0.12	2
	#3	MRSA	0.25	2

PTK796 was also active in vitro against other pathogens isolated as shown in Table 6.

Table 6a. Susceptibility of Other Pathogens (PTK796 Treatment Group)

	#	MIC Range (μg/ml)		
Organism	Isolates	PTK796	Linezolid	Aztreonam
β-hemolytic Streptococcus, Group A	1	0.06	1	-
β-hemolytic Streptococcus, Group B	2	0.06-0.25	1-4	-
β-hemolytic Streptococcus, Group G	2	0.12-0.25	2	-
S. constellatus	1	0.015	2	-
E. faecalis	4	0.06-0.25	2-8	-
S. haemolyticus	1	1	1	-
C. koseri	1	1	-	S
E. cloacae	5	0.06-4	-	S
E. coli	3	0.25-8	-	S
K. pneumoniae	3	2-4	-	S
P. mirabilis	1	8	-	S
A. baumanii	1	0.25		R

* Aztreonam S<4, R>16

Table 6b. Susceptibility of Other Pathogens (Linezolid Treatment Group)

	#	MIC Range (μg/ml)		
Organism	Isolates	PTK796	Linezolid	Aztreonam (S/I/R)
β-hemolytic Streptococcus, Group B	1	0.25	2	-
β-hemolytic Streptococcus, Group G	1	0.25	2	-
S. constellatus	2	0.03-0.06	2	-
E. faecium	1	0.06	2	-
C. freundii	1	8	-	R
E. cloacae	1	2	-	S
E. coli	1	0.5	-	S
K. pneumoniae	2	2-4	-	S

* Aztreonam S<4 R>16

CONCLUSION

The Phase 2 evaluation of PTK796 in comparison to linezolid indicated that PTK was safe and well tolerated and appeared to be efficacious compared to linezolid. The most common pathogen isolated was S. aureus. the majority of which were MRSA. PTK796 was active against S. aureus, with an MIC₈₀ of 0.25µg/ml. Only two clinical failures occurred in the PTK796 treated group. In neither case was the infection caused by an organism with elevated PTK796 MIC. There were three failures in the linezolid treated group and none of these isolates had elevated MICs to linezolid to account for the lack of success.

REFERENCES

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