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Spectrum of Activity of RX-P2382, a Novel Class of Bacterial Ribosome Inhibitor

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Charges may apply.

Introduction

- RX-P2382 is an advanced lead of the pyrrolocytosine class, which has been designed *de novo* to target the ribosome with a novel mechanism of inhibition
 - The structure is shown in Figure 1
- This novel class has been optimized to maximize bacterial influx and minimize efflux to show a broad spectrum of *in vitro* activity across the ESKAPE pathogens
- In this study, RX-P2382's spectrum of activity was examined across a broad range of clinical isolates, including recent drug-resistant pathogens from urinary tract, bloodstream, pneumonia, and other infections
 - Isolates were collected during 2016–2017 from hospitals participating in the global SENTRY Antimicrobial Surveillance Program, 45% of which were from urinary tract infections

Materials and Methods

Figure 1 Structure of RX-P2382



Conclusions

 RX-P2382, a member of the novel pyrrolocytosine class, showed broad-spectrum *in vitro* activity against a wide range of clinical isolates, including Gram-negative and Gram-positive bacteria from various infection types

Acknowledgements

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- RX-P2382 susceptibilities were determined for 594 Gram-negative (GN) and Gram-positive (GP) clinical isolates
 - Isolates tested included 382 Enterobacteriaceae (new taxonomy Enterobacterales; ENT), carbapenem-resistant ENT (CRE), 30
 Pseudomonas aeruginosa (PSA), 31
 Acinetobacter baumannii-calcoaceticus species complex (ACB), 65 Enterococcus spp. including vancomycin-resistant enterococci (VRE), and 32 Staphylococcus aureus including methicillinresistant S. aureus (MRSA)
- JMI Laboratories confirmed all isolate identifications by MALDI-TOF
- Susceptibility testing was performed according to CLSI broth microdilution methodology (M07, M100)
- Isolates were characterized as CRE and extendedspectrum beta-lactamase screen-positive phenotype (ESBL-SP) according to CLSI criteria (M100, 2018)

Results

- RX-P2382 was active against a broad spectrum of clinical isolates, including Gram-negative and Grampositive pathogens, with a unimodal distribution (Tables 1 and 2)
- The MIC_{50/90} values for RX-P2382 were 0.5/1 mg/L against 382 ENT (Table 1)
 - RX-P2382 MIC_{50/90} values were 0.5/4 mg/L, with a range of 0.25 to 8 mg/L for CRE (n = 28)
 - For ESBL-SP isolates, RX-P2382 MIC_{50/90} values were 0.5/1 mg/L, with a range of 0.25 to 8 mg/L (n = 109)

- RX-P2382 was also active against antibioticresistant isolates regardless of resistance mechanism, including ESBL-phenotype, carbapenem-resistance (CRE), vancomycinrestance (VRE) and methicillin-resistance (MRSA)
- The MIC ranges for RX-P2382 were generally narrow and unimodal
- These data suggest that further evaluation of this novel agent is warranted

References

Clinical and Laboratory Standards Institute (2018). M100Ed28E. Performance standards for antimicrobial susceptibility testing: 28th informational supplement. Wayne, PA: CLSI.

Clinical and Laboratory Standards Institute (2018). M07Ed11E. Methods for dilution antimicrobial susceptibility tests for bacteria that grow aerobically; approved standard—eleventh edition. Wayne, PA: CLSI.

Table 1 Antimicrobial activity of RX-P2382 tested against the main organisms and organism groups for Gram-negative species

Organism/organism group (no. of isolates)	No. and cumulative % of isolates inhibited at MIC (mg/L) of ^a :															MIC	
	≤0.004	0.008	0.015	0.03	0.06	0.12	0.25	0.5	1	2	4	8	16	32	> b		
Enterobacteriaceae			-	1		-	1		1	-			1	1			
RX-P2382 (382)					0 0.0	30 7.9	125 40.6	154 80.9	60 96.6	6 98.2	5 99.5	2 100.0				0.5	1
ESBL-phenotype Enterobac	cteriacea	e															
RX-P2382 (109)						0 0.0	39 35.8	50 81.7	13 93.6	1 94.5	4 98.2	2 100.0				0.5	1
Carbapenem-resistant Ente	robacteri	aceae															
RX-P2382 (28)						0 0.0	8 28.6	11 67.9	3 78.6	0 78.6	4 92.9	2 100.0				0.5	4
Acinetobacter baumannii-calc	oaceticus	species	s comple	X	·									•	·		
RX-P2382 (31)						0 0.0	5 16.1	9 45.2	15 93.5	2 100.0						1	1
Carbapenem-resistant Acin	etobacter	r baumai	nnii-calco	baceticu	s specie	s comple	ex		·					·	·		
RX-P2382 (13)							0 0.0	4 30.8	8 92.3	1 100.0						1	1
Pseudomonas aeruginosa																	
RX-P2382 (30)								0 0.0	2 6.7	17 63.3	11 100.0					2	4
Carbapenem-resistant Pseu	udomona	s aerugii	nosa											·	·		
RX-P2382 (5)									0 0.0	2 40.0	3 100.0					4	

^a Intensity of shading is proportional to the number of tested isolates within each row that displays the indicated MIC value. ^b Greater than the highest concentration tested.

- The ACB RX-P2382 MIC_{50/90} values were 1/1 mg/L, with a range of 0.25 to 2 mg/L (n = 31; Table 1)
- PSA RX-P2382 MIC_{50/90} values were 2/4 mg/L, with a range of 1 to 4 mg/L (n = 30; Table 1)
 - 6 PSA isolates were carbapenem resistant and showed RX-P2382 MIC values of 2 to 4 mg/L
- RX-P2382 generally had higher MIC values against Stenotrophomonas maltophilia (MIC_{50/90} 8/>32 mg/L, with a range of 2 to >32 mg/L; n = 32)
- RX-P2382 was also active against Gram-positive pathogens as shown in Table 2
 - For Staphylococcus aureus the $MIC_{50/90}$ were 0.12/0.25 mg/L with a range of 0.06 to 1 mg/L (n = 32) including MRSA ($MIC_{50/90}$ 0.12/0.5; n = 16; Table 2)
 - Against VR *E. faecalis*, RX-P2382 MIC_{50/90} values were 0.12/0.12 mg/L, with a range of 0.06 to 0.12 mg/L (n = 10) and VR *E. faecium* MIC_{50/90} values were 0.25/0.5 mg/L, with a range of 0.06 to 1 mg/L(n = 16; Table 2)
 - RX-P2382 had MIC_{50/90} values of 0.06/0.12 mg/L
 (n = 22; Table 2) against *Streptococcus pneumoniae*

Table 2 Antimicrobial activity of RX-P2392 tested against the main organisms and organism groups for Grampositive species

Organism/organism	No. and cumulative % of isolates inhibited at MIC (mg/L) of ^a :														MIC
group (no. of isolates)	≤0.008	0.015	0.03	0.06	0.12	0.25	0.5	1	2	4	8	16	> b	WIIC 50	
Enterococcus faecalis															
RX-P2382 (33)			0	13	18	2								0.12	0 12
			0.0	39.4	93.9	100.0								0.12	0.12
Vancomycin-nonsusceptible (VRE) Enterococcus faecalis															
RX-P2382 (10)			0	2	8									0.12	0 12
			0.0	20.0	100.0									0.12	0.12
Enterococcus faecium	Enterococcus faecium														
RX-P2382 (32)			0	3	10	15	3	1						0.25	05
			0.0	9.4	40.6	87.5	96.9	100.0						0.20	0.5
Vancomycin-nonsusceptik	Vancomycin-nonsusceptible (VRE) Enterococcus faecium														
RX-P2382 (16)			0	2	4	7	2	1						0.25	05
			0.0	12.5	37.5	81.2	93.8	100.0						0.25	0.5
Staphylococcus aureus															
RX-P2382 (32)			0	8	17	5	1	1						0.12	0.25
			0.0	25.0	78.1	93.8	96.9	100.0						0.12	0.20
Methicillin-resistant S. aureus															
RX-P2382 (16)			0	3	7	4	1	1						0.12	0.5
			0.0	18.8	62.5	87.5	93.8	100.0						0.12	0.5
Streptococcus pneumoniae															
RX-P2382 (22)			0	19	3									0.06	0.12
			0.0	86.4	100.0									0.06	0.12
Penicillin-nonsusceptible	Streptoco	ccus pnei	umoniae			•	•	•	•						
RX-P2382 (11)		-	0	8	3									0.00	0.40
			0.0	72.7	100.0									0.06	0.12

^a Intensity of shading is proportional to the number of tested isolates within each row that displays the indicated MIC value. ^b Greater than the highest concentration tested.