

# In Vitro Activity Evaluation of a Next-Generation Polymyxin, SPR206, against Non-Fermentative Gram-Negative Bacilli Responsible for Human Infections

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## Introduction

- Non-fermentative gram-negative bacilli (NFGNB) are opportunistic organisms that have emerged as important healthcare-associated pathogens, mainly in the immunocompromised patient population
- These organisms are innately resistant to many antimicrobial classes due to the presence of intrinsic genes encoding  $\beta$ -lactamases and decreased permeability
- SPR206 is a polymyxin derivative compound being clinically developed for treating serious infections caused by gram-negative organisms (Figure 1)
- This study evaluated the *in vitro* potency of SPR206 and compared its potency to those of polymyxin-B and colistin against a current collection of NFGNB

## Materials and Methods

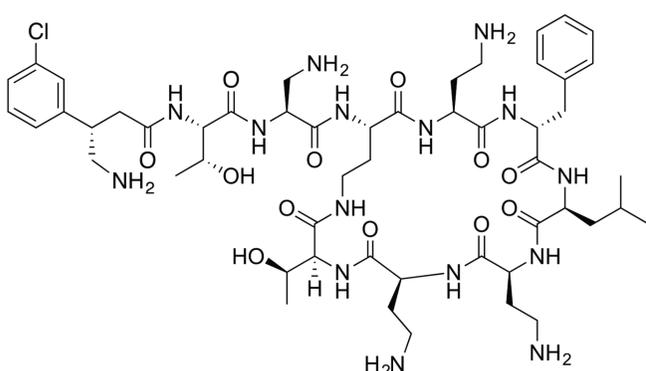
### Bacterial isolates

- A total of 389 randomly selected isolates (182 *Pseudomonas aeruginosa*, 185 *Acinetobacter* spp., and 22 *Stenotrophomonas maltophilia*) representing current antimicrobial susceptibility profiles for the respective species were included
- In addition, a subset of 53 and 130 meropenem-nonsusceptible *P. aeruginosa* and *Acinetobacter* spp., respectively, were analyzed separately
- These isolates originated from the SENTRY Antimicrobial Surveillance Program (2016-2017) bank of organisms and were recovered from 37 medical centers in 20 European nations (n=163), 64 medical centers in the United States (n=203), 12 medical centers from 8 Asia-Pacific nations (n=32), and 12 medical centers from 8 Latin American nations (n=32)
- Isolates were collected from patients with pneumonia (55.4%), skin and skin structure infections (24.7%), bloodstream infections (12.8%), and other infections (7.2%)

### Antimicrobial susceptibility testing

- Isolates were tested against SPR206 and select comparator agents for susceptibility by broth microdilution following guidelines in the Clinical and Laboratory Standards Institute (CLSI) M07 (2018) document
- Frozen-form reference 96-well panels manufactured by JMI Laboratories were used for testing
- Quality assurance was performed by concurrently testing CLSI-recommended quality control reference strains (*Escherichia coli* ATCC 25922, *E. coli* NCTC 13846, *Klebsiella pneumoniae* ATCC 700603, and *P. aeruginosa* ATCC 27853)
- Breakpoint criteria for comparator agents were those available in the CLSI M100 (2018) document

Figure 1 Structure of SPR206



## Results

- SPR206 inhibited all randomly selected *P. aeruginosa* at  $\leq 2$  mg/L and showed MIC results (MIC<sub>50/90</sub>, 0.25/0.5 mg/L) 2-fold lower than colistin (MIC<sub>50/90</sub>, 0.5/1 mg/L) and polymyxin B (MIC<sub>50/90</sub>, 0.5/1 mg/L) (Table 1)
- Similar MIC results for the respective compounds were obtained against carbapenem-nonsusceptible *P. aeruginosa* compared with the randomly selected set (Table 2)
- Against *A. baumannii*, SPR206 (MIC<sub>50/90</sub>, 0.12/0.25 mg/L) was 2- to 8- fold more potent than polymyxin-B (MIC<sub>50/90</sub>, 0.25/1-2 mg/L) and 4- to 32-fold more potent than colistin (MIC<sub>50/90</sub>, 0.5/4-8 mg/L) (Tables 1 and 2)
- In addition, SPR206 inhibited 95.7% of randomly selected *Acinetobacter* spp. or 93.1% of all tested *Acinetobacter* spp. at  $\leq 2$  mg/L
- SPR206 (MIC<sub>50/90</sub>, 0.25/4 mg/L) and polymyxin-B (MIC<sub>50/90</sub>, 0.5/4 mg/L) showed similar MIC values against *S. maltophilia*, and these compounds had MIC results 4- to 16-fold lower than colistin (MIC<sub>50/90</sub>, 4/16 mg/L) (Table 1)

## Conclusions

- Overall, SPR206 showed potent *in vitro* activity against a current collection of NFGNB, and its potency was consistently greater than the clinically available in-class comparator agents
- The *in vitro* results obtained for SPR206 warrant additional investigations to explore its clinical utility for treating infections caused by commonly multidrug-resistant gram-negative pathogens

Table 1 Antimicrobial activity of SPR206 and comparators tested against a random selection of non-fermentative gram-negative bacilli

Organism (no. of isolates)	Antimicrobial agent	Number and cumulative % of isolates inhibited at MIC (mg/L) of <sup>a</sup> :												MIC <sub>50</sub>	MIC <sub>90</sub>	%S <sup>c</sup>			
		$\leq 0.015$	0.03	0.06	0.12	0.25	0.5	1	2	4	8	16	32				64	> <sup>b</sup>	
<i>P. aeruginosa</i> (182)																			
	SPR206			1	19	113	43	5	1								0.25	0.5	
				0.5	11.0	73.1	96.7	99.5	100.0										
	Colistin					15	82	80	3	2							0.5	1	98.9
						8.2	53.3	97.3	98.9	100.0									
	Polymyxin-B					22	139	20	1								0.5	1	100.0
						12.1	88.5	99.5	100.0										
	Meropenem				25	36	35	28	17	8	12				21		0.5	>8	77.5
					13.7	33.5	52.7	68.1	77.5	81.9	88.5				100.0				
<i>Acinetobacter</i> spp. (185)																			
	SPR206				53	90	25	8	1	0	0	1	1				0.12	0.25	
					28.6	77.3	90.8	95.1	95.7	95.7	95.7	96.2	96.8				100.0		
	Colistin					38	78	40	8	4	4	3	3	1	6		0.5	4	88.6
						20.5	62.7	84.3	88.6	90.8	93.0	94.6	96.2	96.8	100.0				
	Polymyxin-B				9	91	35	32	6	2	4	2	3	0	1		0.25	1	93.5
					4.9	54.1	73.0	90.3	93.5	94.6	96.8	97.8	99.5	99.5	100.0				
	Meropenem				1	25	21	11	7	3	4				113		>8	>8	35.1
					0.5	14.1	25.4	31.4	35.1	36.8	38.9				100.0				
<i>S. maltophilia</i> (22)																			
	SPR206			1	8	4	1	1	1	4	1	1					0.25	4	
					4.5	40.9	59.1	63.6	68.2	72.7	90.9	95.5	100.0						
	Colistin				1	0	3	2	4	2	4	4	2				4	16	
					4.5	4.5	18.2	27.3	45.5	54.5	72.7	90.9	100.0						
	Polymyxin-B				2	6	5	1	4	3	1						0.5	4	
					9.1	36.4	59.1	63.6	81.8	95.5	100.0								
	Meropenem														22		>8	>8	
															100.0				

<sup>a</sup> The intensity of shading is proportional to the number of tested isolates within each row that display the indicated MIC value.  
<sup>b</sup> Greater than the highest concentration tested.  
<sup>c</sup> Susceptible breakpoints were those from CLSI/EUCAST (2018).

Table 2 Antimicrobial activity of SPR206 and comparators tested against meropenem-nonsusceptible non-fermentative gram-negative bacilli

Organism (no. of isolates)	Antimicrobial agent	Number and cumulative % of isolates inhibited at MIC (mg/L) of <sup>a</sup> :												MIC <sub>50</sub>	MIC <sub>90</sub>	%S <sup>c</sup>			
		$\leq 0.015$	0.03	0.06	0.12	0.25	0.5	1	2	4	8	16	32				64	> <sup>b</sup>	
<i>P. aeruginosa</i> (53)																			
	SPR206				5	25	19	3	1								0.25	0.5	
					9.4	56.6	92.5	98.1	100.0										
	Colistin					3	27	22	0	0	1						0.5	1	98.1
						5.7	56.6	98.1	98.1	98.1	100.0								
	Polymyxin-B					8	35	9	1								0.5	1	100.0
						15.1	81.1	98.1	100.0										
<i>Acinetobacter</i> spp. (130)																			
	SPR206				40	64	13	4	1	0	0	1	1				0.12	0.25	
					30.8	80.0	90.0	93.1	93.8	93.8	93.8	94.6	95.4				100.0		
	Colistin					22	54	27	6	4	3	3	1	6			0.5	8	83.8
						16.9	58.5	79.2	83.8	86.9	90.0	92.3	94.6	95.4	100.0				
	Polymyxin-B				4	61	22	25	6	2	4	2	3	0	1		0.25	2	90.8
					3.1	50.0	66.9	86.2	90.8	92.3	95.4	96.9	99.2	99.2	100.0				

<sup>a</sup> The intensity of shading is proportional to the number of tested isolates within each row that display the indicated MIC value.  
<sup>b</sup> Greater than the highest concentration tested.  
<sup>c</sup> Susceptible breakpoints were those from CLSI/EUCAST (2018).

## Acknowledgements

This project has been funded in whole or in part with federal funds from the National Institute of Allergy and Infectious Diseases, National Institutes of Health, Department of Health and Human Services, under Contract No. HHSN272201500014C. This study was supported by Spero Therapeutics. JMI Laboratories received compensation for services related to preparing this poster.

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