Monitoring the In Vitro Activity of Tebipenem, an Orally Available **Carbapenem Agent, against a Current Collection of Surveillance Enterobacterales** Clinical Isolates (2018)

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Introduction

- Complicated and uncomplicated urinary tract infections (UTIs) are frequent in any given population
- Current guidelines recommend that empiric antibiotic therapy for UTIs should be based on local resistance data, drug availability, drug intolerance/allergy, and patient history
- Nitrofurantoin or trimethoprim-sulfamethoxazole (TMP-SMX), if local resistance $\leq 20\%$ can be used empirically for uncomplicated cystitis, while fluoroquinolones, ceftriaxone, aminoglycosides, and carbapenems are appropriate for pyelonephritis and complicated UTIs
- However, managing such infections is challenged by the dissemination of isolates producing extended-spectrum β -lactamases (ESBLs) and displaying cross-resistance to fluoroquinolones, aminoglycosides, and folate pathway inhibitors in the hospital and community settings
- Carbapenem agents represent good options for treating these patients with UTIs, who usually have a spectrum of illness severity, comorbidities, and clinical histories, and an oral carbapenem option would provide additional benefits
- Tebipenem, the active metabolite of tebipenem-pivoxil (SPR859), is an oral carbapenem introduced in Japan (2009) for pediatric respiratory and otolaryngologic infections
- SPR994 is a novel tebipenem formulation (tebipenem-pivoxil hydrobromide) that is under clinical development for treatment of UTIs
- This study evaluated the tebipenem activity and spectrum against targeted Enterobacterales species

Materials and Methods

Surveillance isolates

- Enterobacterales isolates included in this study were collected as part of the SENTRY Antimicrobial Surveillance Program
- A total of 1,031 isolates composed of target numbers of randomly selected species recovered from documented infections during 2018 were included (Table 1)
- Preference was given to isolates recovered from urinary tract samples (669 isolates; 64.9%), but those recovered from bloodstream infections (186 isolates; 18.0%) and lower respiratory tract of patients with pneumonia (176 isolates; 17.1%) were included, as well
- Isolates were collected in 113 medical centers located in 26 countries and 9 US census divisions: the United States (674 isolates [65.4%] from 62 medical centers), Europe (262 isolates [25.4%] from 32 medical centers), Latin America (44 isolates [4.3%] from 8 medical centers), and the Asia-Pacific region (51 isolates [4.9%] from 11 medical centers)
- Bacterial species were initially identified by the submitting laboratories and confirmed by JMI Laboratories using standard microbiology methods and matrix-assisted laser desorption ionization-time of flight mass spectrometry (Bruker Daltonics, Bremen, Germany)

Antimicrobial susceptibility testing

- Susceptibility testing was performed for tebipenem and comparator agents by reference broth microdilution method
- 96-well frozen-form broth microdilution panels with cation-adjusted Mueller-Hinton broth (CAMHB) were manufactured by JMI Laboratories (North Liberty, Iowa, USA) per the Clinical and Laboratory Standards Institute (CLSI) specifications described in the M07 (2018) document

Results

- (Figure 2)

Conclusions

- development

Quality control strains were tested before and concomitantly with selected ind bacterial inoculum density was monitored by counting the number of ing units present in the inoculum material

Interpretations of MIC values obtained from clinical isolates used CLSI M100 (2019) breakpoint criteria, as available

The ESBL phenotype was defined for Escherichia coli, Klebsiella pneumoniae, Klebsiella oxytoca, and Proteus mirabilis as an isolate that displayed MIC values $\geq 2 \,\mu g/mL$ for ceftriaxone, ceftazidime, and/or aztreonam

• Overall, tebipenem showed MIC_{50} and MIC_{90} results of 0.06 µg/mL and 0.25 µg/mL, respectively, against all isolates in the study (Table 1)

These tebipenem (MIC_{50/90}, 0.06/0.25 μ g/mL) MIC results were similar to meropenem (MIC_{50/90}, 0.06/0.12 µg/mL)

Isolates displaying tebipenem MIC values of >2 μ g/mL were meropenemnonsusceptible (Table 1)

The lowest tebipenem MIC₅₀ values (0.008 µg/mL) were obtained against Citrobacter koseri and E. coli, followed by Citrobacter freundii, K. pneumoniae, and K. oxytoca $(MIC_{50}, 0.015 \,\mu g/mL)$ (Table 1)

A tebipenem MIC₅₀ value of 0.03 µg/mL was obtained against *Enterobacter cloacae* and Enterobacter aerogenes (Table 1), while the MIC₅₀ value of 0.12 μ g/mL was observed for other species tested (Proteus spp., Morganella morganii, Providencia spp., and Serratia marcescens)

Agents other than carbapenems did not show high activity against *Enterobacterales*, except for piperacillin-tazobactam (91.0% susceptible) and the aminoglycosides (92.5–99.2% susceptible) (Figure 1)

Carbapenems, amikacin, and tigecycline showed susceptibility rates >80% when tested against *Enterobacterales* isolates, meeting the MIC criteria for ESBL phenotype

Tebipenem MIC_{50} values within ± 1 doubling dilution were obtained against the respective species of isolates with and without an ESBL phenotype (Table 2)

Similar MIC₅₀ results were observed for tebipenem when tested against Enterobacterales isolates showing nonsusceptible or susceptible phenotypes for levofloxacin or trimethoprim-sulfamethoxazole (Table 2)

Tebipenem showed broad-spectrum and potent activity against this current and targeted collection of *Enterobacterales* clinical isolates and inhibited all but 14 isolates (98.6%) at $\leq 0.5 \, \mu g/mL$

Tebipenem was most active against E. coli, K. pneumoniae, K. oxytoca, C. koseri, and C. freundii (MIC₅₀, 0.008–0.015 µg/mL) followed by Enterobacter spp., Proteus spp., M. morganii, Providencia spp., and S. marcescens (MIC₅₀, 0.03–0.12 µg/mL)

ESBL phenotype and/or nonsusceptibility to levofloxacin and/or trimethoprimsulfamethoxazole did not adversely affect tebipenem activity

These data suggest that tebipenem may be a convenient oral option for treating UTIs caused by carbapenem-susceptible Enterobacterales isolates and warrant its clinical

Table 1 MIC distributions of tebipen

| Species | No. and cumulative % of isolates inhibited at MIC (μ g/mL) of: | | | | | | | | | | | | | MIO | N/10 |
|-----------------------------|---|-------------|-------------|-------------|-------------|-------------|-------------|------------|-----------|------------|-----------------------|------------|------------|-------|-------|
| (no. of isolates) | ≤0.004 | 0.008 | 0.015 | 0.03 | 0.06 | 0.12 | 0.25 | 0.5 | 1 | 2 | 4 ^a | 8 ª | >8ª | | |
| Enterobacterales (1,031) | 2 0.2 | 113 11.2 | 174 28.0 | 161 43.6 | 195 62.6 | 229 84.8 | 125 96.9 | 18 98.6 | 0 98.6 | 1 98.7 | 3 99.0 | 2 99.2 | 8 100.0 | 0.06 | 0.25 |
| E. coli (102) | 1 1.0 | 67 66.7 | 30 96.1 | 4 100.0 | | | | | | | | | | 0.008 | 0.015 |
| C. koseri (55) | 1 1.8 | 36 67.3 | 17 98.2 | 1 100.0 | | | | | | | | | | 0.008 | 0.015 |
| C. freundii (50) | | 4 8.0 | 32 72.0 | 11 94.0 | 2 98.0 | 0 98.0 | 0 98.0 | 0 98.0 | 0 98.0 | 0 98.0 | 0 98.0 | 1 100.0 | | 0.015 | 0.03 |
| K. pneumoniae (85) | | 1 1.2 | 42 50.6 | 34 90.6 | 4 95.3 | 1 96.5 | 0 96.5 | 0 96.5 | 0 96.5 | 0 96.5 | 0 96.5 | 0 96.5 | 3 100.0 | 0.015 | 0.03 |
| K. oxytoca (16) | | | 10 62.5 | 6 100.0 | | | | | | | | | | 0.015 | 0.03 |
| E. aerogenes (105) | | | 15 14.3 | 44 56.2 | 35 89.5 | 9 98.1 | 2 100.0 | | | | | | | 0.03 | 0.12 |
| E. cloacae (103) | | 5 4.9 | 27 31.1 | 25 55.3 | 26 80.6 | 9 89.3 | 5 94.2 | 0 94.2 | 0 94.2 | 0 94.2 | 2 96.1 | 0 96.1 | 4 100.0 | 0.03 | 0.25 |
| M. morganii (107) | | | | 2 1.9 | 7 8.4 | 45 50.5 | 47 94.4 | 6 100.0 | | | | | | 0.12 | 0.25 |
| P. mirabilis (103) | | | | 5 4.9 | 15 19.4 | 39 57.3 | 38 94.2 | 6 100.0 | | | | | | 0.12 | 0.25 |
| P. rettgeri (49) | | | | 1 2.0 | 19 40.8 | 23 87.8 | 4 95.9 | 1 98.0 | 0 98.0 | 1 100.0 | | | | 0.12 | 0.25 |
| P. stuartii (50) | | | | 2 4.0 | 15 34.0 | 25 84.0 | 8 100.0 | | | | | | | 0.12 | 0.25 |
| S. marcescens (103) | | | 1 1.0 | 8 8.7 | 42 49.5 | 35 83.5 | 10 93.2 | 4 97.1 | 0 97.1 | 0 97.1 | 1 98.1 | 1 99.0 | 1 100.0 | 0.12 | 0.25 |

Meropenem-nonsusceptible.

Figure 1 Antimicrobial activity of comparator agents tested against Enterobacterales and Enterobacterales displaying an extended-spectrum β-lactamase phenotype

| nem tested against surveillance | Enterobacterales | isolates |
|---------------------------------|-------------------------|----------|
|---------------------------------|-------------------------|----------|

Table 2 MIC distributions of tebipenem tested against specific phenotypes

| Phenotypic organism group (no. | No. and cumulative % of isolates inhibited at MIC (μ g/mL) of: | | | | | | | | | | | MIC | | | |
|--|---|------------|-------------|-------------|-------------|-------------|-------------|------------|-----------|-----------|------------|------------|-----------------|----------|-------|
| of isolates) | ≤0.004 | 0.008 | 0.015 | 0.03 | 0.06 | 0.12 | 0.25 | 0.5 | 1 | 2 | 4 a | 8 a | >8 ^a | 50 State | 90 |
| ESBL-phenotype <i>E. coli</i> (20) ^b | | 6 30.0 | 10 80.0 | 4 100.0 | | | | | | | | | | 0.015 | 0.03 |
| Non-ESBL-phenotype <i>E. coli</i> (82) | 1 1.2 | 61 75.6 | 20 100.0 | | | | | | | | | | | 0.008 | 0.015 |
| ESBL-phenotype <i>Klebsiella</i> spp. (25) ^b | | | 12 48.0 | 10 88.0 | 0 88.0 | 0 88.0 | 0 88.0 | 0 88.0 | 0 88.0 | 0 88.0 | 0 88.0 | 0 88.0 | 3 100.0 | 0.03 | >8 |
| ESBL-phenotype carbapenem- susceptible <i>Klebsiella</i> spp. (22) ^b | | | 12 54.5 | 10 100.0 | | | | | | | | | | 0.015 | 0.03 |
| Non-ESBL-phenotype <i>Klebsiella</i> spp. (76) | | 1 1.3 | 40 53.9 | 30 93.4 | 4 98.7 | 1 100.0 | | | | | | | | 0.015 | 0.03 |
| ESBL-phenotype <i>P. mirabilis</i> (7) ^b | | | | 1 14.3 | 1 28.6 | 3 71.4 | 2 100.0 | | | | | | | 0.12 | |
| Non-ESBL-phenotype <i>P. mirabilis</i> (96) | | | | 4 4.2 | 14 18.8 | 36 56.2 | 36 93.8 | 6 100.0 | | | | | | 0.12 | 0.25 |
| Levofloxacin-susceptible Enterobacterales (894)° | 1 0.1 | 93 10.5 | 157 28.1 | 148 44.6 | 179 64.7 | 191 86.0 | 103 97.5 | 16 99.3 | 0 99.3 | 1 99.4 | 1 99.6 | 0 99.6 | 4 100.0 | 0.06 | 0.25 |
| Levofloxacin-non-susceptible Enterobacterales (135) | 1 0.7 | 20 15.6 | 17 28.1 | 13 37.8 | 15 48.9 | 37 76.3 | 22 92.6 | 2 94.1 | 0 94.1 | 0 94.1 | 2 95.6 | 2 97.0 | 4 100.0 | 0.12 | 0.25 |
| Trimethoprim-sulfamethoxazole- susceptible Enterobacterales (855) ^d | 1 0.1 | 88 10.4 | 141 26.9 | 141 43.4 | 180 64.4 | 190 86.7 | 98 98.1 | 14 99.8 | 0 99.8 | 0 99.8 | 1 99.9 | 0 99.9 | 1 100.0 | 0.06 | 0.25 |
| Trimethoprim-sulfamethoxazole- resistant <i>Enterobacterales</i> (176) | 1 0.6 | 25 14.8 | 33 33.5 | 20 44.9 | 15 53.4 | 39 75.6 | 27 90.9 | 4 93.2 | 0 93.2 | 1 93.8 | 2 94.9 | 2 96.0 | 7 100.0 | 0.06 | 0.25 |
| SRI extended-spectrum ß-lactamase | | | | | | | | | | | | | | | |

SBL, extended-spectrum p-lactamase

^c Isolates with MIC results of $\leq 2 \mu g/mL$.

^d Isolates with MIC results of $\leq 2/38 \, \mu g/mL$.



^b The ESBL phenotype was defined for *E. coli, K. pneumoniae, K. oxytoca,* and *P. mirabili*s as isolates that displayed MIC values of ≥2 µg/mL for ceftriaxone, ceftazidime, and/or aztreonam.

Acknowledgements

This study and presentation were sponsored by Spero Therapeutics.

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