

Characterization of Tebipenem Pharmacokinetics-Pharmacodynamics for Efficacy Against Enterobacteriaceae in a One-Compartment *In Vitro* Infection Model

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INTRODUCTION

- Tebipenem (SPR859) is the active form of the orally bioavailable prodrug tebipenem pivoxil (SPR994).
- Tebipenem pivoxil is an oral carbapenem with broad-spectrum *in vitro* activity against Gram-positive, and -negative bacteria and is being developed as an oral option for the treatment of patients with complicated urinary tract infections (cUTI).
- The goal of the studies described herein was to characterize the pharmacokinetics-pharmacodynamics (PK-PD) of tebipenem against a diverse panel of Enterobacteriaceae isolates using a one-compartment *in vitro* infection model. Specific objectives included the following:
 - To carry out dose-fractionation studies to identify the PK-PD index most associated with efficacy of tebipenem against Enterobacteriaceae; and
 - To carry out dose-ranging studies to determine the magnitude of the PK-PD index most associated with efficacy that is required for various levels of bacterial reduction for a panel of Enterobacteriaceae isolates.

METHODS

Antimicrobial Agent and Challenge Isolates

- Tebipenem was provided by Spero Therapeutics (Cambridge, MA).
- A panel of 13 Enterobacteriaceae isolates were supplied from the American Type Culture Collection (ATCC), National Collection of Type Cultures (NCTC) and JMI Laboratories (North Liberty, IA).

In Vitro Susceptibility Testing

- In accordance with Clinical Laboratory Standards Institute (CLSI) guidelines [1], susceptibility studies were completed in triplicate over a two-day period to determine the tebipenem minimum inhibitory concentration (MIC) associated with each Enterobacteriaceae isolate in the challenge panel.

One-Compartment In Vitro Infection Model Dose-Fractionation Studies

- A series of 24-hour dose-fractionation studies were completed using a single *Escherichia coli* isolate (ATCC 25922).
- Bacteria (1 x 10⁸ colony forming units [CFU]/mL) were exposed to tebipenem concentrations that mimicked human healthy volunteer free-drug plasma concentration-time profiles following oral drug administration.
- Seven tebipenem total daily dose levels (24-hour area under the concentration-time curve [free-drug AUC] range, 0.11 to 19.0 mg•h/L) were fractionated in equal divided doses administered every 4, 8, or 12 hours (q4h, q8h, and q12h, respectively).

- Samples were collected for the evaluation of pharmacokinetic (PK) profiles, and enumeration of bacterial burden over the course of the study.

One-Compartment In Vitro Infection Model Dose-Ranging Studies

- In the dose-ranging studies, 13 clinical Enterobacteriaceae isolates were exposed to tebipenem doses ranging from 4.69 to 1200 mg administered q8h (24-hour free-drug AUC ranging from 0.14 to 37.2 mg•h/L).

METHODS

Pharmacokinetic-Pharmacodynamic Analysis

- PK models were fit to the samples collected for the evaluation of the drug concentration profile.
- Data from the dose-fractionation studies were evaluated using Hill-type models and non-linear least squares regression. Relationships between change in log₁₀ CFU/mL from baseline at 24 hours and each of the following tebipenem PK-PD indices were characterized:
 - Free-drug maximum concentration (C_{max}) to MIC ratio (C_{max}:MIC ratio), minimum concentration (C_{min}) to MIC ratio (C_{min}:MIC ratio), percent time above MIC (%T>MIC), and AUC:MIC ratio, with and without adjustment for dosing interval tau for the latter [AUC:MIC ratio•1/tau dosing interval tau [τ]].
- Hill-type models and non-linear least squares regression were also used to evaluate the data from the dose-ranging studies for the PK-PD index most associated with tebipenem efficacy.

RESULTS

In Vitro Susceptibility Testing

- The known resistance mechanisms and tebipenem MIC values of 0.008 to 0.25 mg/L for the isolates evaluated in the *in vitro* infection models are provided in **Table 1**.

Table 1. Tebipenem MIC values and known resistance mechanisms for the isolates evaluated within the one-compartment *in vitro* infection model dose-fractionation and dose-ranging studies

Isolate	Known resistance mechanisms	Tebipenem MIC (mg/L)
<i>E. coli</i> ATCC 25922 ^a	Wild type ATCC reference strain	0.015
<i>E. coli</i> NCTC 13441	CTX-M-15, Sequence Type-131 (ST-131)	0.015
<i>E. coli</i> 4643	CTX-M-15, OXA1/30	0.008
<i>E. coli</i> 13319	CTX-M-15, TEM-1, AcrAB-TolC overexpression OmpC, OmpF porin deficient	0.015
<i>E. coli</i> 872217	CMVII	0.25
<i>E. coli</i> 845741	CTX-M-15, OXA-1, SHV-12, (ST-131)	0.03
<i>Klebsiella pneumoniae</i> 25021	CTX-M-15, TEM-1, OXA-2	0.03
<i>K. pneumoniae</i> 604	CTX-M-15, OXA1/30, SHV-1	0.06
<i>K. pneumoniae</i> ATCC 43816	Wild type ATCC reference strain	0.015
<i>K. pneumoniae</i> 632346	CTX-M-15, OXA-1/30, SHV-5	0.125
<i>K. pneumoniae</i> 934954	CTX-M-15, OXA-1, SHV-28, TEM-1	0.125
<i>P. mirabilis</i> ATCC 29906	Wild type ATCC reference strain	0.015
<i>P. mirabilis</i> ATCC BAA-2791	TEM-1, CMY-4	0.25

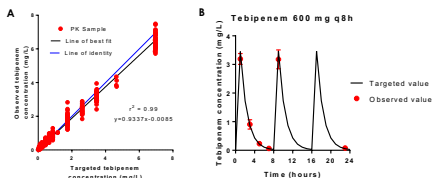
a. Isolate utilized for the dose-fractionation studies.

One-Compartment In Vitro Infection Model Studies

- As evidenced by the agreement between targeted and observed tebipenem concentrations shown in **Figure 1**, the targeted free-drug plasma concentration-time profiles were well simulated for all tebipenem dosing regimens studied in the *in vitro* infection model.

RESULTS

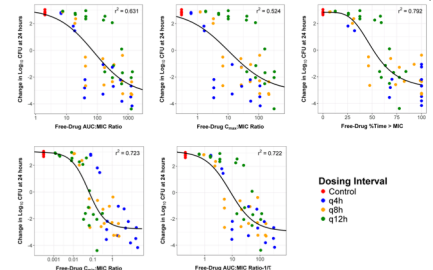
Figure 1. Relationship between all targeted and observed tebipenem concentrations simulated in the one-compartment *in vitro* infection model (A) and the average concentration-time profile for the tebipenem 600 mg q8h regimen (B)



One-Compartment In Vitro Infection Model Dose-Fractionation Studies

- Results of dose-fractionation study data shown in **Figure 2** demonstrated the activity of tebipenem to be time-dependent, with free-drug C_{min}:MIC ratio, %T>MIC and AUC:MIC ratio•1/τ similarly describing the PK-PD of tebipenem based on r² values.
 - Free-drug %T>MIC failed to describe the data well as evidenced by the pooling of data at 100% free-drug %T>MIC, effects ranging from net bacterial stasis to a 4 log₁₀ CFU reduction from baseline.
 - Free-drug C_{min}:MIC ratio failed to describe the *in vitro* activity of tebipenem well as evidenced by the layering of activity by dosing interval, with larger exposures required when tebipenem was administered more frequently.
 - Free-drug AUC:MIC ratio•1/τ was considered to be the PK-PD index most associated with tebipenem efficacy.

Figure 2. Relationships between change in log₁₀ CFU/mL from baseline at 24 hours and each of free-drug tebipenem AUC:MIC ratio, C_{max}:MIC ratio, %T>MIC, C_{min}:MIC ratio, AUC:MIC•1/τ based on dose-fractionation study data

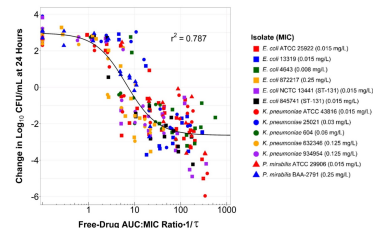


RESULTS

One-Compartment In Vitro Infection Model Dose-Ranging Studies

- The relationship between change in log₁₀ CFU/mL from baseline at 24 hours and tebipenem free-drug AUC:MIC ratio•1/τ based on data from the dose-ranging studies is shown in **Figure 3**.
- The magnitude of free-drug AUC:MIC ratio•1/τ associated with achieving net bacterial stasis and 1- and 2-log₁₀ CFU reductions from baseline based on data for the panel of 13 Enterobacteriaceae isolates was 7.23, 13.1, and 32.4, respectively.

Figure 3. Relationship between the change in log₁₀ CFU/mL from baseline at 24 hours and tebipenem free-drug AUC:MIC ratio•1/τ based on data for 13 Enterobacteriaceae isolates evaluated in the dose-ranging studies



CONCLUSIONS

- The results of dose-fractionation studies demonstrated that free-drug AUC:MIC ratio•1/τ was the PK-PD index most associated with tebipenem efficacy.
- Based upon data for 13 Enterobacteriaceae isolates evaluated in the dose-ranging studies, the magnitude of tebipenem free-drug AUC:MIC ratio•1/τ associated with net bacterial stasis and 1- and 2-log₁₀ CFU reductions from baseline was 7.23, 13.1, and 32.4, respectively.
- These data will be useful to design other pre-clinical studies and support tebipenem dose selection for clinical studies in patients with cUTI.

REFERENCES

- Clinical and Laboratory Standards Institute. Methods for dilution antimicrobial susceptibility tests for bacteria that grow aerobically. Approved standard, 10th edition. CLSI document M07-A10. Clinical and Laboratory Standards Institute, Wayne, PA, 2015.

ACKNOWLEDGEMENTS

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