

Pharmacokinetics of Cefepime-Enmetazobactam in the Cerebrospinal Fluid of Rats

Alexia Chauzy¹, Isabelle Lamarche¹, H el ene Mirfendereski^{1,2}, Ana s Dournayan³, Juan Quevedo⁴, Sandrine Marchand^{1,2}

Contact: alexia.chauzy@univ-poitiers.fr

¹ Universit e de Poitiers, Inserm U1070 PHAR2, Poitiers, France

² CHU de Poitiers, service de Toxicologie-Pharmacocin etique, Poitiers, France

³ ADVANZ PHARMA SAS, Paris, France

⁴ ADVANZ PHARMA S.L.U, Madrid, Spain

Introduction

Nosocomial meningitis caused by Gram-negative pathogens remains a major therapeutic challenge, particularly with the increasing prevalence of multidrug-resistant bacteria that restrict available treatment options. Cefepime is standard care, but its activity is increasingly compromised by β -lactamase-mediated resistance [1]. Enmetazobactam, a novel extended-spectrum β -lactamase inhibitor, has been developed to restore the antibacterial spectrum of cefepime. To evaluate the potential applicability of this combination for central nervous system (CNS) infections, we investigated the pharmacokinetics of cefepime/enmetazobactam in cerebrospinal fluid (CSF) of rats.

Materials and methods

- Experiments were conducted in accordance with EC Directive 2010/63/EU after approval by the local ethics committee (authorization number: 55247-2025042216508773).
- Thirty-eight healthy Sprague-Dawley rats weighing 308 ± 67 g were used.
- Cefepime/enmetazobactam was administered as an intravenous bolus via the tail vein at a dose of 50/25 mg/kg in anesthetized rats.
- Blood and CSF samples were collected over 210 min in a minimum of 3 animals per sampling time. Blood samples were collected by intracardiac puncture while CSF was collected from the cisterna magna. Plasma was separated from whole blood after centrifugation at 11000 g for 10 min at 4 C.
- Cefepime/enmetazobactam concentrations in plasma and CSF were determined by LC-MS/MS. The plasma protein binding of both compounds was found to be negligible in prior investigations and was consequently assumed to be zero [2].
- A sparse non-compartmental pharmacokinetic analysis was conducted using PKNCA R package [3].

Results

- Mean PK parameters are presented in Table 1.
- Compared with plasma, CSF concentration-time profiles were flat for both compounds (Figure 1), reflecting slow and limited distribution, with CSF-to-plasma AUC ratios of 4.93% and 4.86% for cefepime and enmetazobactam, respectively.

Table 1. Mean pharmacokinetic parameters obtained by sparse noncompartmental analysis in healthy rats after intravenous bolus administration of 50/25 mg/kg of cefepime (FEP)/enmetazobactam (META).

Matrix	Parameter	Unit	FEP	META
Plasma	C ₀	mg/L	284	314
	t _{1/2}	min	25.2	22.3
	AUC _{0-∞}	h*mg/L	109	85
	CL	L/(h*kg)	0.460	0.294
	V _d	L/kg	0.279	0.157
CSF	C _{max}	mg/L	1.59	1.57
	T _{max}	min	60	20
	t _{1/2}	min	108	85.3
	AUC _{0-∞}	h*mg/L	5.36	4.16

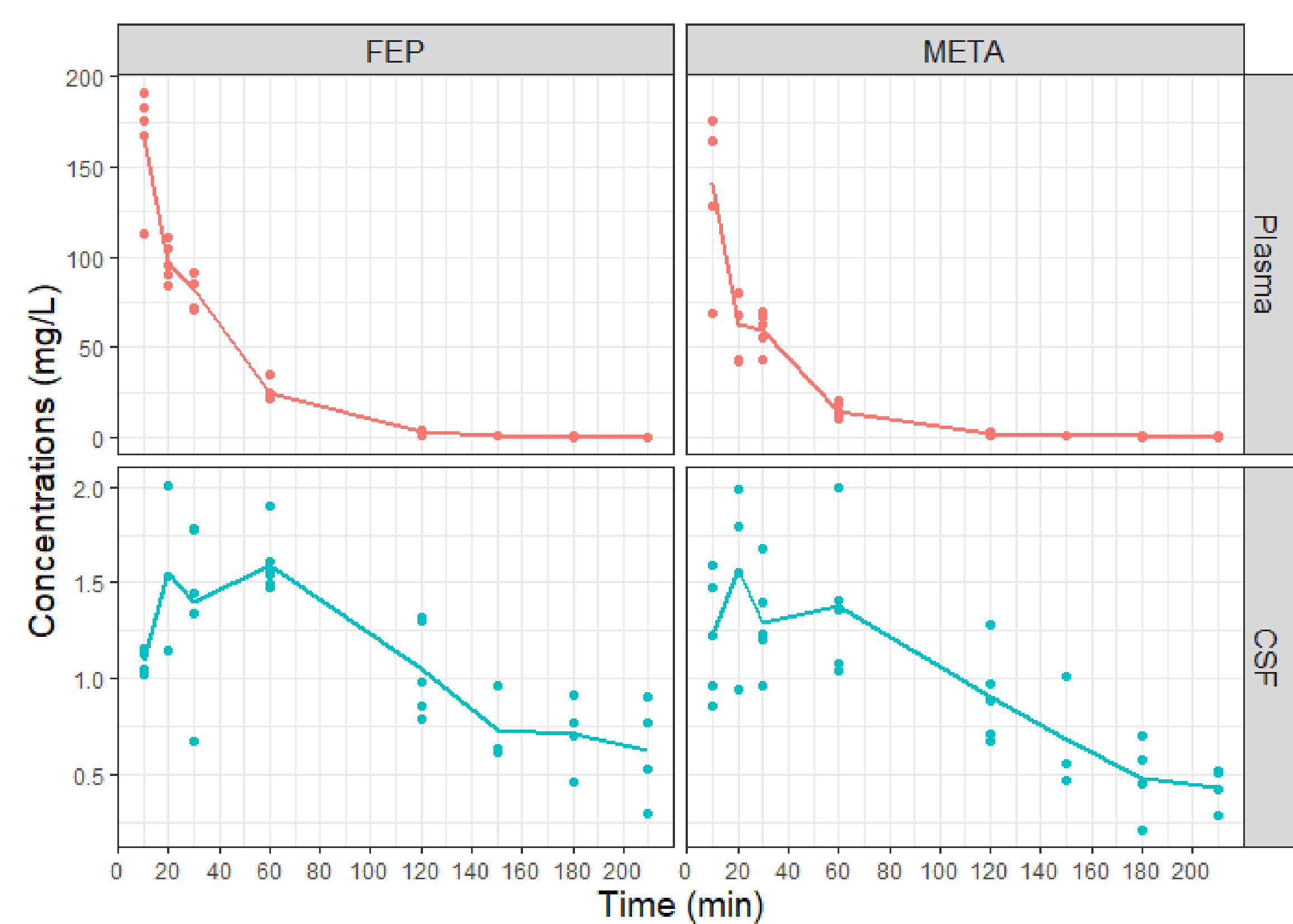


Figure 1: Plasma and CSF concentrations of cefepime (FEP) and enmetazobactam (META) in rats (n=3-5 per time point).

Conclusion

Although both compounds showed limited CSF penetration, the cefepime AUC ratio was consistent with the range of previously reported values [4]. Importantly, these results provide the first available CSF penetration data for enmetazobactam. Overall, our findings provide initial pharmacokinetic data to further investigate the potential role of cefepime/enmetazobactam in CNS infections.

References:

[1] Gorham et al. 10.1080/14787210.2025.2565581

[3] Denney et al. 10.1007/s10928-015-9432-2

[2] Crandon et al. 10.1128/AAC.00033-15

[4] Lodise et al. 10.1016/j.diagmicrobio.2005.09.007

Funding: This project benefits from financial support from ADVANZ PHARMA Services (UK) Ltd

