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 Simple and accurate quantitative analysis of cefiderocol and ceftobiprole in human plasma using liquid chromatography-isotope dilution tandem mass spectrometry:

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Abstract

37 Objectives: Cefiderocol and **ceftobiprole** are new generation cephalosporin antibiotics that exhibit high inter-individual plasma concentration variability that potentially impact their efficacy or toxicity. The aim of this study was to develop and validate a selective, simple and 40 fast UPLC-MS/MS method for simultaneous quantification of cefiderocol and **ceftobiprole** in 41 human plasma to enable their therapeutic drug monitoring and support PK and PK/PD studies, in particular in critically ill patients.

 Methods: After a simple and fast single-step protein precipitation, cefiderocol and 44 ceftobiprole were separated on a Waters Acquity UPLC BEH C18 column by linear gradient elution; with subsequent detection by Shimadzu MS 8060 triple quadrupole tandem mass spectrometer in a positive ionization mode.

 Results: Analysis time was 5 minutes per run. The analytical performance of the method in terms of specificity, sensitivity, linearity, precision, accuracy, matrix effect, extraction recovery, limit of quantification, dilution integrity and stability of analytes under different conditions met all criteria for a bioanalytical method for the quantification of drugs. The 51 calibration curves were linear over the range of 1-200 mg/L for cefiderocol and 0.5-100 mg/L 52 for **ceftobiprole** with a linear regression coefficient above 0.995 for both.

 Conclusion: A simple, fast, and selective liquid chromatography-tandem mass spectrometry method was developed and validated for the simultaneous quantification of cefiderocol and 55 ceftobiprole. This new method was successfully applied to the measurement of plasma 56 concentration of cefiderocol and **ceftobiprole** in critically ill patients and showed good performance for their therapeutic monitoring and optimizing antibiotic therapy.

58 Keywords: liquid chromatography, mass spectrometry, cefiderocol, ceftobiprole, cephalosporin, therapeutic drug monitoring.

 against a large range of bacteria involved in pneumonia, including Gram-positive bacteria (GPB), such as methicillin-resistant *Staphylococcus aureus* (MRSA), *Enterococcus faecalis*, and *Streptococcus pneumoniae*, and Gram-negative bacteria (GNB), such as *Pseudomonas aeruginosa* and *Haemophilus influenza* [1]. Hospital-acquired pneumonia (HAP) and community-acquired pneumonia (CAP) are among the most common infections treated in the hospital setting [2, 3]. Two randomized, double-blind, phase III clinical trials have 68 demonstrated the efficacy of **ceftobiprole** in patients with HAP or CAP [4, 5]. Ceftobiprole exhibits high inter-individual pharmacokinetic (PK) variability [6]. Moreover, an exposure- efficacy relationship [7] and an exposure-toxicity relationship [8] have been demonstrated. The PK variability and the pharmacokinetics/pharmacodynamics (PK/PD) relationship of 72 ceftobiprole suggest the potential interest of its routine therapeutic drug monitoring (TDM). Cefiderocol (S-649266) is a newly *Food and drug administration* (FDA) and *European Medicines Agency* (EMA) approved, first in its class, parenterally administered siderophore cephalosporin with a novel mechanism to penetrate the outer cell membrane of GNB, including multidrug-resistant strains [9]. The emergence of carbapenem resistance in many GNB, involved in a variety of serious infections including pneumonia, urinary tract infections, intra-abdominal infections and bloodstream infections, is an urgent threat to global public health [10-13], and the presence of multi-drug resistance complicates the management of 80 these infections due to the limited treatment options available [14, 15]. Data from global surveillance studies for cefiderocol have shown potent *in vitro* activity against a wide range of GNB, including carbapenem-resistant *Acinetobacter baumannii*, *Pseudomonas aeruginosa*,

Ceftobiprole, a fifth-generation parenteral cephalosporin, has shown antimicrobial activity

Enterobacteriaceae and *Stenotrophomonas maltophilia* strains [16]. The noninferiority and

tolerability of cefiderocol versus imipenem-cilastatin was demonstrated in a randomized,

85 double blind, phase II study for the treatment of patients with complicated urinary tract 86 infections due to multidrug-resistant GNB [17]. The PK profile of cefiderocol has been described in phase I and II single- and multiple-dose clinical studies [18-20]. PK and safety have also been described in subjects with renal impairment [19, 20]. However, the population PK and PK/PD properties of cefiderocol in ill patients, and specifically in critically ill patients are currently limited [21]. This is particularly important because there are, in this population, several co-morbidities and modifications of the physiological state which can modify the PK and PK/PD profiles of most beta-lactam antibiotics used in clinical practice [22-26].

 The aim of this study was to develop and validate a selective, simple and fast UPLC-MS/MS 94 method for simultaneous quantification of cefiderocol and **ceftobiprole** in human plasma to enable their TDM and support PK and PK/PD studies.

- **2. Materials and methods**
-

2.1. Chemical and reagents

99 Cefiderocol and $[2H_{12}]$ -cefiderocol powders were provided by Shionogi & Co, Ltd (Osaka, 100 Japan) while **ceftobiprole** (BAL9141, the active form) and $[2H₄]$ -ceftobiprole powders were provided by Basilea Pharmaceutica Ltd (Basel, Switzerland). The chemical structures and mass spectrum of cefiderocol and ceftobiprole are shown in **Figure 1** (mass spectrum of [2H12]-cefiderocol and [2H4]-ceftobiprole are shown in **Supplemental Figure 1**). Methanol was obtained from Merck (Darmstadt, Germany). Formic acid and ammonium acetate were obtained from Sigma-Aldrich (Munich, Germany). Sulphosalicylic acid dihydrate (SSA) and acetonitrile were obtained from VWR (Fontenay-sous-Bois, France). All reagents used were of the highest available analytical grades. Liquid chromatography–MS/MS grade water (ultra-pure water) was purchased from a water distribution hypergrade system Purelab Flex®

 (ELGA®), and drug-free plasma from healthy donors was supplied by the French Blood Establishment (Paris, France).

2.2. Preparation of stock solutions, standards and quality control samples

112 Individual stock solutions of cefiderocol and **ceftobiprole** were prepared at 1,000 mg/L. 113 Cefiderocol was prepared in ultra-pure water while **ceftobiprole** was prepared in a solution of 114 methanol-hydrochloric acid 1N (95:5, v/v). Working solutions of cefiderocol and ceftobiprole, obtained by diluting the stock solution with methanol, were used to spike drug-free plasma to prepare independent calibration standards and quality control (QC) samples. Calibration ranges covered in the current method are: 1-200 mg/L (1-5-10-25-50-100-200) for cefiderocol 118 and $0.5-100 \text{ mg/L } (0.5-1-5-10-25-50-100)$ for **ceftobiprole**. The QC samples were tested at four different concentrations: high QC (HQC: 80% of upper limit of quantification [ULOQ]), medium QC (MQC: 50% of selected range), low QC (LQC: 2 times the lower limit of 121 quantification [LLOQ]) and QC at LLOQ. A solution of mix (ISmix) of $[2H_{12}]$ -cefiderocol 122 and $[2H_4]$ -ceftobiprole, used as internal standard, at 10 mg/L for both, was prepared in ultra- pure water. Stock solutions, working solutions, calibration standards, ISmix and QC samples were stored at -80°C.

2.3. Instruments and analytical conditions

 Chromatography was performed on a Nexera X2 system (Shimadzu, Japan) with an autosampler temperature at 8°C. Acquity UPLC BEH C18 column (4.6 x 50 mm, 3.5 µm particle size) was used for chromatographic separation and column temperature was 129 maintained at 45°C. Separation of cefiderocol and **ceftobiprole** was achieved over 5 min using two mobile phases: (A) water-formic acid (100:0.1, v/v)-ammonium acetate 2 mM and (B) acetonitrile-formic acid (100:0.1, v/v). A binary pump delivered the mobile phases at a flow rate of 0.5 mL/min using a linear gradient elution. The UPLC system was coupled to a triple quadripole mass spectrometer: MS 8060 (Shimadzu, Japan). Quantifications were achieved in Multiple Reactions Monitoring (MRM) mode and electrospray ionization (ESI) was operated 135 in positive mode. The interface temperature, the desolvatation line (DL) temperature and the heat block temperature were set at 300°C, 250°C and 400°C, respectively, with a drying gas flow of 10.0 L/min and a nebulizing gas flow of 3.0 L/min. The interface voltage was set at 4.0 kV. Argon was used as collision gas. Chromatographic data acquisition; peak integration and quantification were performed using LabSolutions Insight LC-MS 3.2 SP1 software.

2.4. Samples pre-treatment

141 Sample preparation was performed by protein precipitation: 10 μ L of 30% sulphosalicylic acid and 50 µL of ISmix were added to 100 µL of human plasma, calibrator or QC samples. The mixture was vortexed for 1 min and centrifuged for 10 min at 18,900 g at room temperature. Subsequently, 5 µL of the clear supernatant was transferred to an autosampler 145 vial containing 200 μ L of mobile phase A. 1 μ L was injected into the LC-MS/MS system.

2.5. Method validation

 The validation was performed according to *European Medicines Agency* (EMA) guideline for the validation of bioanalytical methods [27]. Parameters included were selectivity, carry-over, linearity, accuracy and precision, lower limit of quantification, matrix effect, stability in human plasma and dilution integrity.

151 2.5.1. Selectivity

152 The analytical method should be able to differentiate the analyte(s) of interest from 153 endogenous components in the matrix or other components in the sample such as other drugs. 154 To perform this, fifteen different sources of plasma samples were tested. A selective method 155 should not have interference of more than 20% of the LLOQ for the analytes and 5% for the ISs.

2.5.2. Carry-over

158 Carry-over was assessed by injecting drug-free plasma samples after a high concentration 159 calibrator. Carry-over in the drug-free plasma samples following the high concentration calibrator should not be greater than 20% of the LLOQ.

2.5.3. Linearity

 Calibration curve was acquired by plotting the peak area ratio of the concentration of 163 cefiderocol and **ceftobiprole** standards to the area of their respective isotopic IS over the range 164 from 1 to 200 mg/L for cefiderocol and 0.5 to 100 mg/L for **ceftobiprole**. Curves were 165 assayed by least square weighted $(1/x)$. Linearity was defined by a linear regression 166 coefficient $(R)^2 \ge 0.995$.

2.5.4. Precision and accuracy

 The intra-day precision and accuracy were evaluated using six different replicates, extracted in the same day, at the four QC levels (LLOQ, LQC, MQC and HQC). The inter-day precision and accuracy were determined by repeating each QC levels six times a day for three 171 consecutive days ($n = 18$ replicates). The concentration of each QC levels was determined using calibration standards prepared on the same day. The precision was calculated as the coefficient of variation (CV, %) within a single run (intra-day assay) and between different runs (inter-day assay), and the accuracy as the percentage ratio of the measured and nominal 175 concentration (mean of measured/nominal \times 100). The acceptance limits were CV <15% for 176 precision and within \pm 15% of the nominal concentration for accuracy (ranged from 85-115%), except for the LLOQ at which deviation of 20% is acceptable.

2.5.5. Lower limit of quantification

2.5.6. Matrix effect and extraction recovery

 Matrix effect (ME) and extraction recovery (ER) were assessed at two QC levels (LQC and HQC) in quintuplicate with five different sources of plasma. The approach involves determination of ratio of peak areas of analytes in three different sets [28]. One consisting of 188 analyte standards in methanol (set A), one prepared in drug-free matrix extracts and spiked 189 after extraction (set B), and one prepared in drug-free matrix from the same sources but spiked before extraction (set C). ME and ER were calculated by the following equations: ME $(\%) = B/A \times 100$ and ER $(\%) = C/B \times 100$. Normalized ME was obtained by dividing the 192 values reported for the analytes by those of the IS. A value above or below 100% for the ME indicates an ionization enhancement or suppression respectively. The CV obtained for the IS-normalized ME should be less than 15%.

2.5.7. Stability

196 The stability in human plasma of cefiderocol and **ceftobiprole** in different storage conditions was performed at LQC, MQC and HQC. During each day tested, we used freshly prepared 198 calibrators and each QC levels were analysed on three different replicates. The value was estimated by comparing measured concentration before storage to after storage, and reported as the percentage ratio of mean measured concentration to nominal concentration. The short term stability at room temperature (bench-top stability) was evaluated by measuring each QC level, stored at 25°C, every hour for 8 consecutive hours. The short term stability in the fridge was determined after 24h at 4-8°C and the short term stability in the freezer was determined

204 after 1 week at -25°C. The long term stability was determined after 2 month storage at -80°C. 205 The freeze and thaw stability was estimated after three complete freeze and thaw cycles of - 206 80°C to 25°C. Stability of extracts kept onboard the autosampler at 8°C during 24h was also 207 tested. Cefiderocol and **ceftobiprole** was considered to be stable in plasma, during this 208 different storage conditions, when measured concentration within \pm 15% of the nominal 209 concentration.

210 2.5.8. Dilution integrity

 The dilution integrity was examined to ascertain that an unknown sample with concentration 212 exceeding the upper limit of compound calibration range, could be diluted with drug-free matrix without influencing the accuracy and precision of the measurement. To achieve this, a sample was prepared in a two-fold higher concentration than of the ULOQ (400 mg/L for 215 cefiderocol and 200 mg/L for **ceftobiprole**) followed by dilution (1:3) in drug-free plasma before extraction. Diluted sample was tested for accuracy and precision: the acceptance limits 217 were CV <15% for precision and within \pm 15% of the nominal concentration for accuracy. Samples were analyzed in quintuplicate.

219 2.6. Clinical application

220 This UPLC-MS/MS quantification method was applied to measurement of cefiderocol and 221 ceftobiprole in plasma of critically ill patients hospitalized in ICU at Pitié-Salpêtrière hospital 222 (Paris, France) to perform TDM and PK studies. Ten patients with compassionate use of 223 cefiderocol and twenty five patients treated with **ceftobiprole** were included in the study. 224 Cefiderocol and **ceftobiprole** were administrated by intravenous infusion over 3-hours and 225 over 2-hours, respectively, and two successive blood samples were collected into lithium 226 heparin tubes at steady state for both: one prior to the start of the infusion (C_{trough}) and one 15 227 ± 10 min after the end of the infusion (C_{max}). Plasma samples were prepared by centrifuging collected blood samples for 5 min at 4,500 *g* at room temperature. All plasma samples were frozen at -80°C until analysis, and were process and analysed as previously described. French regulations on non-interventional observational studies do not require patient's consent when analyzing data obtained from routine care. Approval for data collection was obtained from the Commission Nationale de l'Informatique et des Libertés (n°1491960v0).

3. Results

3.1. Optimization of LC-MS/MS conditions

 Electrospray positive mode yielded a better spectrometer response than the negative mode. To achieve symmetrical peak shapes, good resolution and a short chromatographic run time, a 239 mobile phase consisting of (A) water-formic acid (100:0.1, v/v)-ammonium acetate 2 mM and 240 (B) acetonitrile-formic acid (100:0.1, v/v) was used in the experiments using gradient elution: the gradient was initiated at 3% B, then increasing linearly to 51.3% B in 2 min, after which it was directly changed to 100% B, where it was maintained for 1 min, before returning to the initial condition for the last 2 min, resulting in a total run time of 5 min. Mass spectrometry 244 parameters for the LC-MS/MS determination of cefiderocol, **ceftobiprole** and their respective IS are shown in **Table 1**.

3.2. Method validation

3.2.1. Selectivity and carry-over

248 Fifteen different sources of **heparin-plasma samples** without analytes but containing the following anti-infective drugs: levofloxacin, ciprofloxacin, moxifloxacin, rifampicin, ceftazidime, cefepime, amoxicillin, voriconazole, posaconazole, fluconazole and isavuvonazole were tested. These anti-infective drugs were tested due to their relatively common use in ICU and for their possible concomitant administration in our cohort of patients. No interference with endogenous compounds or tested anti-infective drugs was 254 observed above 20% of the LLOQ of cefiderocol or **ceftobiprole** and above 5% of their

255 respective IS. Furthermore, these same fifteen heparin-plasma samples were spiked at the

256 LLOQ of cefiderocol and ceftobiprole and were tested for accuracy and precision No

- 257 significant interference was found in spiked plasma samples compared to non-spiked samples.
- 258 The accuracy (% true) and precision (% CV) of the spiked samples were: cefiderocol (112;
- 259 10.7) and ceftobiprole (109; 11.2) indicating an absence of significant interference that could
- 260 influence the accuracy and precision of the measurement.
- 261 The carry-over observed was less than 20% of the LLOQ for both drugs [cefiderocol (8.4%);

262 ceftobiprole (1.8%)]. Furthermore, no carry over was observed for the IS used. The MRM-

263 chromatogram of drug-free plasma sample of ceftobiprole and cefiderocol are shown in

- 264 **Figure 2A4 and 2B4** respectively.
- 265 3.2.2. Linearity

266 Calibration curve were linear with linear regression coefficient $(R)^2 \ge 0.995$ for both analytes 267 (0.998 and 0.999 for cefiderocol and **ceftobiprole** respectively). All calibrators, analyzed on 268 seven different days, were measured with an accuracy ranged from 87-111% and coefficient 269 of variation less than 10.2%. The highest calibration point was defined as ULOQ. The MRM-270 chromatogram of ceftobiprole and cefiderocol at their ULOQ are shown in **Figure 2A2 and** 271 **2B2** respectively. The linear regression equations of pooled data obtained over seven distinct 272 days were: $y = 0.20360x - 0.08026$ and $y = 0.72831x - 0.07829$ for cefiderocol and 273 ceftobiprole respectively.

274 3.2.3. Accuracy and precision

275 Intra- and inter-day precision and accuracy outcomes of QC samples are shown in **Table 2**. 276 The intra- and inter-day coefficients of variation ranged from 2.6% to 5.3% and from 4.5% to 277 10.4% respectively, for both analyte at all tested concentrations (LQC, MQC and HQC).

3.2.4. Lower limit of quantification

 The LLOQ was established at 0.5 mg/L for ceftobiprole and at 1 mg/L for cefiderocol (**Table 2**). The MRM-chromatogram of **ceftobiprole** and cefiderocol at their LLOQ are shown in **Figure 2A1 and 2B1** respectively.

3.2.5. Matrix effect and extraction recovery

 Table 3 presents the matrix effect (absolute and IS-normalized) and extraction recovery data 286 at low and high concentrations using heparin-plasma samples containing different anti-287 infective drugs (see list in selectivity section). For cefiderocol, a significant ion enhancement was observed, but this matrix effect was compensated by the use of its isotope-labeled IS. The CV of the IS-normalized matrix effect was less than 15% for both analyte. The absolute extraction recovery for both analyte ranged from 90 to 103%, with in all instances CVs below 10%.

3.2.6. Stability

 Analyte stability was determined at five different storage conditions, using three concentrations of QC (LQC, MQC, HQC), as percent ratio of mean measured concentration to nominal concentration. Results are summarized in **Table 4**. The long term stability proved that both analytes are stable in plasma for at least 2 months at -80°C and after three cycles of freeze and thaw. The short term stability proved that both analytes are stable for 8h at room 298 temperature; 24h at 4 $\textdegree C$; for at least use at -25 $\textdegree C$ and for 24h in the autosampler after 299 extraction. In addition, the stock, working and ISmix solutions have also been stable at $-80^{\circ}C$ for at least 2 months.

 The accuracy (% true) and precision (% CV) of the diluted sample were: cefiderocol (94; 5.7) 303 and c eftobiprole (106; 7.2).

3.3. Clinical application

 This validated UPLC-MS/MS method was successfully applied to the measurement of plasma 306 concentration of cefiderocol and **ceftobiprole** in critically ill patients. For cefiderocol, the 307 usual dosing was $2g$ TID [three times a day] (6/10). Three patients with renal impairment 308 were treated with $0.75g$ BID [twice a day], 1g TID and 1.5g TID regimen, respectively, and 309 one patient with renal hyperfiltration was treated with 2g QID [four times a day] regimen. In 310 all patients, cefiderocol was infused over 3-hours. Concerning the **ceftobiprole**, all patients were treated with 500mg TID regimen infused over 2-hours. Both analytes were easily detected and measured in patients' plasma. The results are represented in **Figure 3**. Moreover, as shown in **Figure 2A5 and 2B5**, no interferences were observed between ceftobiprole or cefiderocol and endogenous compounds or others drugs given to participating ICU patients.

4. Discussion

 We present here a liquid chromatography-isotope dilution tandem mass spectrometry method 318 for the simultaneous measurement of cefiderocol and **ceftobiprole**. Based on the high recovery, relatively low intra- and inter-day CVs, and good linearity, the present method is suitable for detection and quantification of these two antibiotics in human plasma. To the best of our knowledge, very few analytical methods intended for routine cefiderocol or ceftobiprole clinical application have yet been published. Lima *et al.* reported an HPLC-UV 323 method for measurement of ceftobiprole in human plasma with concentration range of 1-80 mg/L [8]. Likewise, Mernissi *et al.* and Zimmer *et al.* reported an HPLC-UV method for

325 routine therapeutic drug monitoring of cefiderocol with concentration range of 2.5-100 mg/L 326 and 4-160 mg/L, respectively [29, 30]. Although HPLC-UV has the advantage of being cost-327 effective and easily accessible to most laboratories, HPLC-MS/MS method operating the 328 multiple reaction monitoring has become one the top choices for high speed selective and 329 sensitive analysis of compounds including drugs [31]. HPLC-MS/MS has been employed for 330 cefiderocol and **ceftobiprole** PK studies in healthy volunteers [18, 32, 33] but in these reports, 331 information about the analytical methods is lacking making their reproducibility difficult. Our 332 method is the first method allowing the simultaneous determination of cefiderocol and 333 ceftobiprole by mass spectrometric detection (analysed using their respective isotopic internal 334 standard) and designed to perform their therapeutic monitoring in a routine setting.

335 The specificity of our method was acceptable since we did not observed interference with 336 endogenous compounds or drugs given to patients hospitalized in ICU. Likewise, the 337 sensitivity was sufficient with a first calibration point (1 mg/L for cefiderocol and 0.5 mg/L 338 for **ceftobiprole**) well below the mean trough concentration measured in clinical studies and in 339 our patients. This new method uses a wide concentration range which makes it suitable for the 340 measurements of concentration for future pharmacokinetic studies. In comparison to other 341 methods intended for routine clinical application, the current one has an extended lower and 342 upper limit of quantification for both antibiotics. Given the potentially high plasma 343 concentration variability, this could be interesting in case of very low concentration at C_{min} 344 and very high concentration at C_{max} . In addition, compared to the previously reported HPLC-345 UV methods for which the run times analysis varies from 12-15 min [8, 29, 30], our method is 346 rapid with fast sample preparation and run times (5 min) and requires only a small volume of 347 plasma (100µL), which could reduce the time required for quantification of large number of 348 samples and the blood volume collected from the patients.

349 Concerning stability, we found that cefiderocol and **ceftobiprole** samples should be dispatched

350 to the laboratory without delay due to their poor stability at room temperature \langle <24h). This is an important information for the management of the pre-analytical stage and to manage shipping of samples coming from other hospitals. Likewise, both antibiotics were stable for at least 2 months at -80°C in plasma. This is particularly important for clinical research protocols, where samples may need to be stored for a long time before they can be assayed. Our stability results for cefiderocol were consistent with those reported by Mernissi *et al*. [29], but significantly differs from those reported by Zimmer *et al*. [30] in particular for 357 bench-top and fridge stability.

 Concerning the clinical application, conducted in critically ill patients, our results suggest that there is significant inter-individual plasma concentration variability; this argues to consider 360 cefiderocol and **ceftobiprole** as candidates for TDM in this patient population. A PK/PD studies should be performed to explore if this exposure variability can influence their efficacy or toxicity and to define their therapeutic targets concentration.

5. Conclusion

 We have developed and validated a rapid, sensitive, selective, accurate, precise and reliable 366 UPLC-MS/MS method for the simultaneous quantification of cefiderocol and ceftobiprole in human plasma. This method was successfully applied to their therapeutic monitoring and will allow us to carry out PK/PD studies in critically ill patients.

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 Informed consent: French regulations on non-interventional observational studies do not require patient's consent when analyzing data obtained from routine care.

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B

Molecular Formula: $C_{20}H_{22}N_8O_6S_2$
Formula Weight: 534.56868 Formula Weight:
[M+H]+: **CEFTOBIPROL**

[M+H]+: 535.117647 Da

Figure 1. Chemical structures and mass spectrum of (A) cefiderocol and (B) ceftobiprole

ACeftobiprole

BCefiderocol

Q 752.10>285.20 (+) 8.24e3

Figure 2. MRM ion-chromatograms of (A1) blank human plasma spiked with ceftobiprole at 0.5 mg/L (LLOQ), (A2) blank human plasma spiked with ceftobiprole at 100 mg/L (ULOQ), (A3) d4-ceftobiprole (IS of ceftobiprole) at 10 mg/L, (A4) blank human plasma without ceftobiprole, (A5) real patient sample treated with ceftobiprole at Ctrough, (B1) blank human plasma spiked with cefiderocol at 1 mg/L (LLOQ), (B2) blank human plasma spiked with cefiderocol at 200 mg/L (ULOQ), (B3) d12-cefiderocol (IS of cefiderocol) at 10 mg/L, (B4) blank human plasma without cefiderocol, (B5) real patient sample treated with cefiderocol at 15min after the end of infusion.

Figure 3. Observed (A) cefiderocol (n = 10) and (B) ceftobiprol (n = 25) plasma concentration at steady-state Ctrough (the median time of blood collection was 2 [1-5] days after start treatment) and at Cmax (samples were collected at 15 \pm 10 min after the end of infusion). The box plot show the interquartile and extreme range, the mean (+) and median (horizontal line) values for each group.

A. [2H12]-cefiderocol

B. [2H4]-ceftobiprole

Supplementary Figure 1. Mass spectrum of (A) $[2H_{12}]$ -cefiderocol and (B) $[2H_4]$ -ceftobiprole

Table 1. Analyte quantification characteristics

CE: collision **energy**

Table 2. Summarized results of the precision and accuracy experiments.

NC: nominal concentration, n: number of replicate, CV: coefficient of variation

Table 3. Absolute matrix effect (ME), IS-normalized ME and absolute extraction recovery (ER) at low and high concentration with their corresponding CVs (n = 5).

n: number of replicates

n: number of replicates