

**Manuscript** Ferrone V. et al. Meropenem, levofloxacin and linezolid in human plasma of critical care patients: A fast semi-automated micro-extraction by packed sorbent UHPLC-PDA method for their simultaneous determination. Pharm Biomed Anal. 2017;140:266-273. doi: 10.1016/j.jpba.2017.03.035.

**Title** Meropenem, levofloxacin and linezolid in human plasma of critical care patients: a fast semi-automated micro-extraction by packed sorbent UHPLC-PDA method for their simultaneous determination

**Article type** Full length article

### Abstract

An ultra high-performance liquid chromatographic (UHPLC) method with PDA detection was developed and validated for the simultaneous quantification of meropenem, linezolid, and levofloxacin in human plasma and applied in human plasma of critical care patients. A semi-automated microextraction by packed sorbent (MEPS) for sample preparation was used. All parameters in the extraction step (pH, sample volume, sample dilution and number of aspiration - ejection cycles) and in the desorption step (percentage of acetonitrile in the solvent of elution and number of aspirations of elution solvent through the device) were statistically significant when the recovery was used as response. The method showed good linearity with correlation coefficients,  $r^2 > 0.9991$  for the three drugs, as well as high precision (RSD% < 10.83% in each case). Accuracy ranged from -7.8% to +6.7%. The limit of quantification of the three drugs was established at 0.01  $\mu\text{g/mL}$  for linezolid and levofloxacin and 0.02  $\mu\text{g/mL}$  for meropenem. Linezolid, meropenem, levofloxacin and the internal standard were extracted from human plasma with a mean recovery ranged from 92.4% to 97.4%. During validation, the concentration of meropenem, linezolid and levofloxacin was found to be stable after 3 freeze-thaw cycles and for at least 24 h after extraction. This method will be subsequently used to quantify the drugs in patients to establish if the dosage regimen given is sufficient to eradicate the infection at the target site.

**Keywords** meropenem; linezolid ; levofloxacin; MEPS; UHPLC; critical care patients

**Taxonomy** High-Performance Liquid Chromatography With Sub-3-Micron Particle, Sample Extraction, Therapeutic Drug Monitoring

**Manuscript category** Bioanalytical Applications

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Dear Editor,

We are sending our manuscript entitled “Meropenem, levofloxacin and linezolid in human plasma of critical care patients: a fast semi-automated micro-extraction by packed sorbent UHPLC-PDA method for their simultaneous determination” to your journal after all the authors have seen and approved the final form of it. This paper reports the development of an analytical method with a novel sampling and clean-up procedure of biological samples using the micro extraction with packed sorbent (MEPS). At the best of our knowledge there is no MEPS-UHPLC-PDA method in literature for the simultaneous determination of meropenem levofloxacin and linezolid in human plasma.

We declare that we have not published it anywhere else before nor is it being considered simultaneously in another journal and we have no conflict of interest to declare. Also, it is an honour to publication of my paper in your excellent journal.

The following address can be used for any kind of negotiations.

[gcarlucci@unich.it](mailto:gcarlucci@unich.it)

We do look forward to hearing from you soon.

Sincerely yours

Prof. Giuseppe Carlucci

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## Response to Editor

Dear Editor

The authors thank the referees and the editor, for your excellent revision of our paper. In complying with your remarks we have answered point-by-point all their queries. Please find enclosed our responses.

*Combined comments from the reviewers:*

*-In the guide for authors the following restriction is reported: "In the case of bioanalytical (pharmacokinetic, bioequivalence) studies, too general, oversimplified sentences such as "The methods in the literature are not sensitive enough, require tedious sample preparation steps; the retention times are too long and the method requires expensive instrumentation and special skill." are not acceptable." This means that the sentence reported at lines 84-86 should be cancelled, or rephrased.*

Thanks for the comment. The authors agree with the referees, according with their comment the sentences reported at line 84-86 have been cancelled.

*-Furthermore, searching in the literature was not accurate, i.e. the following paper reporting the simultaneous bioanalysis of antibiotics (including those considered in the present papers) by UHPLC-MS/MS was not referenced:*

*Simultaneous analysis of antibiotics in biological samples by ultra high performance liquid chromatography-tandem mass spectrometry, Cazorla-Reyes R., Romero-Gonzalez R., Frenich A.G., Rodriguez Maresca M.A., Martinez Vidal J.L. (2014) Journal of Pharmaceutical and Biomedical Analysis, 89, pp. 203-212.*

The authors thanks the referees for their suggestion. Two new references for the analysis of antibiotics (including those considered in the present papers) by UHPLC-MS/MS have been added.

*-Line 108: regimes ??*

The authors apologize for the typo. Regimens has been added instead of regimes

*-Lines 110-111: the sentence was not grammatically correct and should be amended.*

The authors thank for the comment. The sentence has been corrected.

### *Experimental*

*-Line 145: the core-shell technology of the column should be emphasized.*

Thanks for the insightful comment. Core shell technology has been emphasized in the result and discussion section.

*-Line 150: One full stop was enough.*

The authors apologize for the typo. According with the referees the sentence has been corrected

*-Lines 177 and 195: g should be reported instead of rpm.*

According with the referee g units has been used instead of rpm

*-A section dealing with the real sample analysis and reporting the administered dose of the drugs, the blood volume collected, etc., was missing.*

Thanks for the comment. A new section dealing with real sample analysis has been added.

### *Results and discussion*

*-Line 247: Fused-Core® is a trademark of Advanced Materials Technology, Inc., otherwise, core-shell could be used instead.*

Thanks for the comment. Core-shell has been used instead of fused-core

*-Section 3.3 was a bit confusing. It should report only the validation issues. A new section dedicated to the results and discussion of real samples analysis should be added.*

*The authors agree with the referees. Section 3.3 has been edited according with the comments. Furthermore, a new section dedicated to the results and discussion of real sample analysis has been added.*

### *Figures*

*-A chromatogram at the LOQ should be presented in Fig. 5.*

*A chromatogram at the LOQ has been added to Fig.5 according with the comment.*

We do look forward to hearing from you soon.

Sincerely yours

Prof. Giuseppe Carlucci

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1 **Meropenem, levofloxacin and linezolid in human plasma of critical**  
2 **care patients: a fast semi-automated micro-extraction by packed**  
3 **sorbent UHPLC-PDA method for their simultaneous determination**  
4

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29 **Abstract**

30 An ultra high-performance liquid chromatographic (UHPLC) method with PDA detection was developed and  
31 validated for the simultaneous quantification of meropenem, linezolid, and levofloxacin in human plasma  
32 and applied in human plasma of critical care patients. A semi-automated microextraction by packed  
33 sorbent (MEPS) for sample preparation was used. All parameters in the extraction step (pH, sample volume,  
34 sample dilution and number of aspiration - ejection cycles) and in the desorption step (percentage of  
35 acetonitrile in the solvent of elution and number of aspirations of elution solvent through the device) were  
36 statistically significant when the recovery was used as response. The method showed good linearity with  
37 correlation coefficients,  $r^2 > 0.9991$  for the three drugs, as well as high precision (RSD% < 10.83% in each  
38 case). Accuracy ranged from -7.8% to +6.7%. The limit of quantification of the three drugs was established  
39 at 0.01 µg/mL for linezolid and levofloxacin and 0.02 µg/mL for meropenem. Linezolid, meropenem,  
40 levofloxacin and the internal standard were extracted from human plasma with a mean recovery ranged  
41 from 92.4% to 97.4%. During validation, the concentration of meropenem, linezolid and levofloxacin was  
42 found to be stable after 3 freeze-thaw cycles and for at least 24 h after extraction. This method will be  
43 subsequently used to quantify the drugs in patients to establish if the dosage regimen given is sufficient to  
44 eradicate the infection at the target site.

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46

47

48 **Keywords:** meropenem, linezolid, levofloxacin, MEPS, UHPLC, human plasma, critical care patients.

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## 52 1. Introduction

53

54 The growing drug resistance of *Gram-positive* (e.g. *Staphylococcus Aureus*) and *Gram-negative*  
55 (e.g. *P. Aeruginosa*) has become an issue of concern for clinicians. In presence of growing  
56 resistance, how to optimize the better efficacy or shorter course and how to minimize the  
57 emergence of drug resistance and reduce adverse reactions are issues worthy of further discussion  
58 [1-4]. Combinatory regimes of linezolid with fluoroquinolones (levofloxacin) or beta-lactam  
59 antibiotics (meropenem) are frequently used for initial treatment of severe multidrug-resistant  
60 nosocomial infections, because they provide considerably large broad and partly complementary  
61 antibacterial spectra.

62 Meropenem (MERO), or (4*R*,5*S*,6*S*)- 3-[[*(3S*,5*S*)-5-[(Dimethylamino)carbonyl]-3-pyrrolidinyl]thio]-6-  
63 [[*(1R*)-1-hydroxyethyl]-4-methyl-7-oxo-1-azabicyclo[3.2.0.]hept-2-ene-2-carboxylic acid (Fig.1A), is  
64 a semi synthetic antibiotic member of a class of  $\beta$ -lactam antibiotics, the carbapenems, having a  
65 broad-spectrum and a concentration-dependant bactericidal activity against Gram-positive and  
66 Gram-negative respiratory pathogens. Unlike imipenem, meropenem proves a good stability  
67 against human renal dehydropeptidase I and does not require the coadministration of a  
68 dehydropeptidase enzyme inhibitor [5].

69 Lizenolid (LINZ) or N-[[*(5S*)-3-[3-Fluoro-4-(4-morpholinyl)phenyl] -2-oxo-5-oxazolidinyl] methyl]  
70 acetamide (Fig.1B) is an oxazolidinone derivative with a predominantly bacteriostatic effect  
71 against *Gram-positive* bacteria, including methicillin-resistant *Staphylococcus aureus* (MRSA),  
72 vancomycin-resistant enterococci and cephalosporin-resistant *Streptococcus pneumonia* [6].

73 Levofloxacin (LEVO) or (-)-*(S)*-9-fluoro-2,3-dihydro-3-methyl-10-(4-methyl-1-piperazinyl)-7-oxo-  
74 7*H*-pyrido[1,2,3-*de*]-1,4-benzoxazine-6-carboxylic acid (Fig.1C) is a fluoroquinolone antibiotic  
75 which is a concentration-dependant antibacterial agent with its therapeutic results closely related  
76 to the ratio of the area under the concentration time curve to the minimum inhibitor  
77 concentration (AUC/MIC) for the organism. The quantification of the plasma concentrations for  
78 MERO, LINZ and LEVO could be an important pharmacological tool in order to optimize the drug  
79 dosage regimens and to maximize the ratio of efficacy to resources. Various analytical methods  
80 used for the determination of MERO [7-10] LEVO [11-13] and LINZ [14-17] concentration in  
81 biological matrices have been reported. High-performance liquid chromatography (HPLC) coupled  
82 with photodiode array detector (PDA) [7-9,11] and fluorescence [13] detection are the most

83 widely used techniques. Recently, HPLC coupled with mass spectrometry (LC-MS/MS) for  
84 determination MERO,LEVO or LINZ in human plasma have also been published [18,19].

85 A selective and sensitive ultra-high performance liquid chromatography (UHPLC-PDA) methods are  
86 preferable to more expensive LC-MS techniques. Furthermore, most of the proposed methods are  
87 still quite time-consuming and laborious, requiring large amount of sample and often also the use  
88 of toxic organic solvents in order to achieve satisfactory analytical limits. Clearly, this macro-  
89 approach to sample preparation is not sensitive enough when low volumes of sample are  
90 available. It is well known that recent trends in the field of sample preparation have been focused  
91 on miniaturization, automation, high-throughput performance, on-line coupling with analytical  
92 instruments and cost-effectiveness through extremely low or no solvent consumption in order to  
93 reduce laboratory workload and increase analytical performance. In such settings, micro  
94 extraction by packed sorbent (MEPS) is an attractive option and powerful sample-preparation  
95 approach suitable for accomplishing analytical and bioanalytical challenges. MEPS is a miniaturized  
96 form of the solid-phase extraction (SPE) technique. In MEPS, compared with SPE, the sorbent  
97 material (about 2 - 4 mg) is integrated directly into the syringe and not in a separate column and  
98 can be connected on-line to gas chromatography (GC), liquid chromatography (LC), LC coupled to  
99 mass spectrometry (LC-MS) or GC-MS [20-23].

100 Many sorbent materials, such as silica and silica-based (C<sub>2</sub>, C<sub>8</sub>, C<sub>18</sub>), strong cation exchangers (SCX),  
101 polymeric HDVB and SDVB or molecular imprinted polymers (MIPs) can be used as packing bed or  
102 as a coating. The MEPS technique has been used to extract a wide range of analytes from  
103 biological sample, such as plasma, dialyzed, urine and blood [24-28]. The current research study  
104 describes the development and validation of a sensitive, fast, efficient, reliable and high  
105 throughput semiautomatic MEPS-based methodology in combination with UHPLC-PDA for the  
106 simultaneous determination of MERO, LINZ and LEVO in human plasma to establish if the dosage  
107 regimens given are sufficient to eradicate the infection at the target site. This method was  
108 validated according to the FDA [29] and ICH [30] guidelines. At present, no papers have been  
109 published for the simultaneous analysis of MERO. LEVO, LINZ in human plasma sample using  
110 MEPS-UHPLC-PDA

111

## 112 **2. Materials and methods**

113

## 114 2.1 Chemicals and reagents

115 Meropenem trihydrate (CAS 119478-56-7), linezolid, (CAS 165800-03-3) and levofloxacin (CAS  
116 100986-85-4) were purchased from Sigma-Aldrich (Sigma-Aldrich, Milan, Italy), while ulifloxacin  
117 (CAS 112984-60-8), used as the internal standard (IS) to compensate for variability in extraction  
118 efficiency, was supplied from Suzhou Bichal Biological Technology (Jiangsu, China). Hydrochloric  
119 acid triethylamine and ammonium acetate were obtained from Carlo Erba Reagents (Carlo Erba,  
120 Milan, Italy). Methanol, acetonitrile HPLC grade, were purchased from VWR International (VWR  
121 International, Milan, Italy). All chemicals were of analytical-reagent grade or better. Pooled drug-  
122 free human plasma was obtained from Sigma-Aldrich (Sigma-Aldrich, Milan, Italy).

123

## 124 2.2 Instrumentation

125

126 The analysis of MERO, LINZ, LEVO and the IS, was performed on a Waters Ultra Performance Liquid  
127 Chromatography (ACQUITY H-Class) with column heater, degassing system combined with a  
128 Acquity quaternary solvent manager equipped with Acquity UPLC sample manager and a Waters  
129 2996 PDA detector. Empower v.3 software (Waters) was used for setting-up the analysis and for  
130 data management. A XS104 Mettler Toledo analytical balance was used to weigh the analytes for  
131 the preparation of stock solutions and calibration standards. Water HPLC-grade water obtained by  
132 passage through an Elix 3 and Milli-Q Academic water purification system (Millipore, Bedford, MA,  
133 USA). Extraction was performed using the MEPS 250  $\mu$ L syringe and the MEPS BIN (barrel insert  
134 and needle) containing 4 mg of silica C<sub>18</sub> sorbent from SGE Analytical Science (LabService Analytica,  
135 Italy). Pooled drug-free human plasma was obtained from Sigma-Aldrich (Sigma-Aldrich, Milan,  
136 Italy). A CMA 100 microinjection pump (CMA Microdialysis AB, Kista, Sweden) was used for  
137 semiautomatic control of the flow-rate during aspiration and elution. All solvent evaporations  
138 were carried out in a Visiprep Vacuum Manifold equipped with a Visidry Drying Attachment  
139 System (Supelco, Bellefonte, USA). Samples were sonicated in an ultrasonic bath Labsonic (FALC,  
140 Milan, Italy).

141

## 142 2.3 Analytical and chromatographic conditions

143

144 Chromatographic separation was achieved using a core shell Ascentis Express C<sub>18</sub> (50 x 2.1 mm I.D.  
145 2.7 µm particle size) column protected by a disposable Guard Cartridge Ascentis Express C<sub>18</sub> (5.0 x  
146 2.1 mm) (Sigma-Aldrich, Milan, Italy) maintained at 25±1°C, using a thermostatically controlled  
147 column heater. Mobile phase consisted in 10 mM ammonium acetate buffer adjusted to pH 5.5  
148 with hydrochloric acid (phase A) and a mixture of acetonitrile and methanol in a ratio of 80/20  
149 (v/v) (phase B), both phases were added with 0.1% (v/v) of triethylamine. To perform the best  
150 separation of the analytes a linear gradient elution program was used. Starting from 97% and 3%  
151 of phase A and B, respectively, within 2.5 min the composition of the eluting mixture was turned  
152 to 65% and 35% of phase A and B, respectively, then an isocratic plateau was programmed for 0.8  
153 min and returned back to the original ratio between 3.5 and 3.6 min, followed by 1.5 min of re-  
154 equilibration of the column to the initial condition. The flow rate was set at 0.6 mL/min. The  
155 solvents were filtered before use through a 0.45 µm PTFE membrane (Waters, Milan, Italy), while  
156 ammonium acetate solution was filtered through a 0.45 µm GH-Polypro membrane (VWR  
157 International, Milan, Italy). For quantification purposes, the UV detection was conducted using the  
158 maximum absorbance wavelength of each analyte as follows: 300 nm for meropenem, 254 nm for  
159 Linezolid, 290 nm for levofloxacin and 272 for the IS. Under these conditions, the total run time  
160 was 5 min. The system suitability test (SST) was performed under optimised chromatographic  
161 conditions. It was evaluated as the reproducibility of retention time and area expressed as RSD%.  
162 The results are shown in Table 1.

163

## 164 2.4 Preparation of calibration standard and quality controls

165

166 Stock solutions of MERO, LINZ, LEVO and the IS at the concentration of 2.0 mg/mL were  
167 individually prepared by dissolving 20 mg of each reference powder into 10 mL volumetric flask  
168 with a mixture of methanol and hydrochloric acid 0.1M (95/5 v/v) and stored at -20°C; under these  
169 conditions they were stable for over 4 months. The working solutions were prepared by dilution of  
170 the stock solutions (calibration and quality control) in water. A working solution of I.S. (2.5 µg/mL)  
171 was prepared in water. Standards for the calibration curves and quality control samples were then

172 prepared using serial dilutions of stock solution in drug-free human plasma. A seven-point  
173 calibration curve was built by spiking the blank plasma with appropriate aliquots of the working  
174 solutions to obtain final concentration in the range 0.02 to 30.0 µg/mL for MERO, 0.01 to 10.0  
175 µg/mL for LEVO and 0.01 to 30.0 for LINZ, respectively. 50 µL of plasma sample were diluted to  
176 150 µL with water, sonicated for 10 min at room temperature, then centrifuged (4320 g, 10 min,  
177 4°C). The MEPS needle was fitted with a barrel insert and needle assembly (BIN) containing a C<sub>18</sub>  
178 sorbent. The CMA/100 microinjection pump is a digitally-controlled dispensing unit used to draw  
179 and discharge samples and solutions through the BIN. Before being used for the first time, the  
180 sorbent was conditioned firstly with 250 µL of methanol and then with a mixture water-methanol  
181 (95/5, v/v). This step activates the sorbent and ensures reproducible retention of the analytes.  
182 Aliquots of 50 µL of spiked samples and standards were diluted to 150 µL with water and then  
183 passed through the C<sub>18</sub> sorbent 10 times at a flow rate of 7.5 µL/sec. The sorbent was then washed  
184 with 100 µL of a mixture of 100 mM ammonium acetate buffer (pH 5.5) and methanol (95/5, v/v)  
185 to remove interferences, at a speed of 7.5 µL/sec. Then the analytes were eluted with 150µL of  
186 methanol at a flow rate of 10.0 µL/sec. Pooled quality control (QC) samples of MERO, LINZ and  
187 LEVO were prepared independently in the same way described above at concentration levels  
188 representing the lower limit of quantification (LLOQ), the low (LQC), middle (MQC) and high (HQC)  
189 ranges of the calibration curves to determine the intra- and inter-assay precision and accuracy of  
190 the method. In all measurements (standards and samples) 20 µL of ulifloxacin (giving a 2.5 µg/mL  
191 concentration) were added as internal standard (IS) prior to extraction.

192

## 193 2.5 Sample collection

194 Blood samples were collected at "S.S. *Annunziata*" Hospital (Chieti, Italy) 4h after an intravenous  
195 administration of MERO (Merrem 1g three times a day), LEVO (Levoxacin 400mg twice daily) and  
196 LINZ (Zyvox 600mg twice daily). Sample were collected at the same time as routine TDM samples  
197 in glass tube containing EDTA as the anticoagulant and did not require extra visits to the clinic or  
198 additional fingerpricks. Plasma samples were obtained after centrifugation at 1800 g for 10 min at  
199 4°C, then, the supernatant (plasma) was stored at -20°C. Samples were thawed just before the  
200 extraction procedure and subjected to the method described above.

201

## 202 **3. Results and discussion**

203

### 204 *3.1 Optimisation of MEPS extraction procedure*

205

206 MEPS presents many more optimization opportunities to be investigated. Several parameters  
207 namely, sorbent type, sample dilution, number of extraction cycles, loading volume, sampling flow  
208 rate, elution solvent and its volume, elution flow rate and solvent for conditioning and washing  
209 step .The sorbents used are obviously a critical parameter and the correct choice of the sorbent  
210 will define the analytical performance of the whole methodology. So far, firstly several sorbent  
211 materials, such as silica and silica-based (C<sub>2</sub>, C<sub>8</sub>, C<sub>18</sub>), strong cation exchangers (SCX), polymeric  
212 HDVB and SDVB were investigated in order to evaluate which had the best recoveries, C<sub>18</sub>  
213 achieved best recoveries as shown in Fig 2.

214 Samples can be loaded directly or diluted if they are too concentrated or viscous in order to avoid  
215 the BIN saturation and the consequently the limitation on MEPS extraction efficiency. Two plasma  
216 dilutions with Milli-Q water (1:3, 1:2, v/v) and undiluted plasma were tested. Plasma diluted with  
217 Milli-Q water 1:2 (v/v) was the optimum in terms of reproducibility and sensitivity, undiluted  
218 plasma was unsuitable due to the short sorbent life and the higher irreproducibility of the MEPS  
219 extraction while when plasma diluted 1:3 (v/v) was investigated, lower sensitivity was observed.

220 The sample volume used was investigated in order to obtain the best equilibrium between good  
221 analytical performance and good extraction yield in the 50-250 µL range. In MEPS the analytes  
222 preconcentration on the sorbent phase is affected by the number of extraction cycles performed.  
223 According to Rani et al. (2012), the analytes extraction increased as the number of extraction  
224 cycles enhanced up to ten [31]. Fig 3 shows the results obtained. 150 µL was found to be a good  
225 choice as it was an appropriate volume for all evaluated dilutions. So far, plasma diluted 1:2 and a  
226 sampling volume of 150 µL could be set as optimal conditions. Sampling flow rate was also  
227 evaluated within 1-20 µL/s in order to investigate and modulate the interaction of target analytes  
228 to the sorbent. The optimum was achieved using 10 µL/s, higher values promoted great signal  
229 irreproducibility for the majority of tested analytes (Fig.4). The elution capacity of different  
230 solvents with different polarity was evaluated in term of peak area of each analyte, better results  
231 were obtained when methanol was used; consequently, latter one was selected as the elution  
232 solvent (150 µL ). Different washing solutions were evaluated, including neat ultrapure water and

233 ultrapure water containing methanol at different proportions (5, 10, 25 and 50% v:v). In all cases,  
234 100  $\mu$ L of the washing solution was passed through the cartridge. Best results were obtained when  
235 100  $\mu$ L of a mixture of 100 mM ammonium acetate buffer (pH 4.0) and methanol (95/5 v/v) was  
236 used due to the highest analytical signal of the analytes and the lowest interference of co-eluting  
237 analytes. In order to avoid or reduce problems related to carryover, 6 wash-discard cycles (3 with  
238 100  $\mu$ L methanol and 3 with 100  $\mu$ L water were implemented after elution). This cleaning  
239 procedure removed most of the residual analytes, as the carryover checked after the washing  
240 procedure was reduced to less than 0.5% peak area of the initial amount of analytes in all cases.  
241 The sample clean-up by MEPS is very quick, and additional operations between sample  
242 preparation and injection were not required. The small amount of stationary phase can be easily  
243 and effectively cleaned, avoiding carry-over effect so a single BIN can be used up to 80 times  
244 without loss in extraction performance.

245

### 246 *3.2 Optimization of the chromatographic conditions.*

247 Chromatographic separations were tried by using different compositions of mobile phases at  
248 different pH, different ratios and different analytical columns in order to achieve good resolution,  
249 selectivity and efficiency in a short run time. Higher separation efficiency was obtained using  
250 Ascentis express C<sub>18</sub>. This column, packed with core shell particles, avoids the band broadening  
251 caused by multiple inter-particle paths, reduces the longitudinal diffusion and allows to work at  
252 higher flow rates with lower working pressures compared to totally porous columns. Different  
253 organic modifiers e.g. methanol and acetonitrile were tried in isocratic mode firstly then in  
254 gradient elution. Due to the different chromatographic properties of LINZ, MERO and LEV, isocratic  
255 separation was found to be unsuitable. The optimal mobile phase consisted of a mixture of  
256 methanol-acetonitrile and 10 mM ammonium acetate buffer (pH 4.0) in a gradient elution. Several  
257 gradient elution were explored, best separation was obtained using the gradient elution reported  
258 in Section 2.3. Furthermore to increase the peak shape, different triethylamine percentage in the  
259 range of 0.05 and 0.5 % (v/v) was added to the mobile phase. The optimum was achieved by using  
260 0.1% (v/v) of triethylamine. Under this condition, MERO, LEVO and LINZ, were well separated and  
261 their peak shape were symmetric. Each drug was detected at its maximum wavelength in order to  
262 obtain higher sensitivity. By applying the chromatographic condition herein reported, the total run  
263 time in this assay was 3 min and the retention times for MERO, LEVO and LINZ were 1.55( $\pm$ 0.03),

264 2.36( $\pm$ 0.02) and 2.82( $\pm$ 0.02) min respectively while for the IS was 2.58( $\pm$ 0.02) min with  
265 consistently excellent reproducibility of less than 1.3%. All the evaluate parameters were within  
266 the required range, as established in FDA guidelines [29,30].

267

### 268 3.3 Method validation

269 In order to demonstrate the suitability of the developed analytical method, validation was carried  
270 out according to the International Guidelines [29,30]. Selectivity of the proposed method was  
271 assessed by analyzing six different batches of control blank samples. The absence of any signal at  
272 the same retention time of the selected antibiotics suggested that there were no matrix  
273 interferences that may give false positive. The intra- and inter-day precision (RSD values) and  
274 precision (BIAS %), obtained from the analysis of three batches of LLOQ, QCL, QCM and QCH, in  
275 duplicate, on the same day and for five consecutive days, did not exceed 10.83 % for precision and  
276 ranged between -7.8 and +6.7 for accuracy, as shown in Table 2.

277 The LOQs of the method were 0.02  $\mu$ g/mL for MERO and 0.01  $\mu$ g/mL for LINZ and LEVO,  
278 respectively, with good precision and accuracy. The LODs were 7.0 ng/mL for MERO and 3.0 ng/mL  
279 for LEVO and LINZ. Results are shown in Table 3.

280 Calibration graphs were obtained by a linear regression analysis with weighting factors consisting  
281 in  $1/x^2$  values in order to describe the relationships between concentration of the analytes and  
282 their peak area ratio to I.S. as analytical response. Quantitative analyses for MERO, LEVO and LINZ  
283 were performed at different wavelengths, as reported in Table 1, in order to obtain higher  
284 sensitivity and better signal-to-noise ratio.

285 Carry-over was investigated by injecting into the UHPLC two extracted blank plasma spiked with  
286 MERO, LEVO and LINZ at the ULOQ concentration, followed by three blank samples. No significant  
287 carry over effect (< 0.5%) was evident.

288 Three dilutions (1:4, 1:3 and 1:2) with the blank matrix were evaluated, with at least five  
289 determinations for each dilution factor in order to demonstrate the dilution integrity. For all  
290 evaluated dilutions accuracy was within  $\pm$  8%, while precision, expressed as relative standard  
291 deviations, was less than 6%. The extraction recovery of MERO, LEVO and LINZ from plasma was  
292 determined by assaying aliquots of drug-free plasma samples spiked at the QCL, QCM and QCH  
293 concentrations, results are reported in Table 4. Stability of MERO, LEVO and LINZ was investigated  
294 during sample collection, after long-term storage and short-term storage, and through several



295 freeze and thaw cycles according to the guidance for industry on the validation of bioanalytical  
296 methods [29,30]. The stock solution was stable at room temperature for 48 h and at 4 °C for two  
297 weeks. MERO, LEVO and LINZ were also stable up to 24 h at room temperature in plasma; it  
298 remained intact at -20 °C for up to five weeks. No degradation was observed after three cycles of  
299 freezing and thawing. The stability of MERO, LEVO and LINZ in the extracts was confirmed after a  
300 24 h storage at 4 °C.

301

### 302 *3.4 Application of the proposed method*

303

304 The proposed method was applied for the determination of MERO,LEVO and LINZ in human  
305 plasma samples from patients 4h after being treated with an intravenous administration of MERO  
306 (Merrem 1g three times a day), LEVO (Levoxacin 400mg twice daily) and LINZ (Zyvox 600mg twice  
307 daily). To ensure quality results, an internal quality control was performed in each batch of  
308 samples, which implies a blank sample, a matrix matched calibration and a spiked blank samples at  
309 7.5 µg/mL, 2.5µg/mL and 15µg/mL for MERO, LEVO and LINZ, respectively (Fig.5). The results  
310 obtained are shown in table 5.

311

## 312 **4.Conclusion**

313

314 The determination of drugs in biological matrix is the keystone in drug discovery and drug  
315 development as well as for pharmacodynamic and pharmacokinetic studies. Patients attending  
316 hospital emergency services with severe multi-drug resistant infections need rapid dosage  
317 adjustment in order to eradicate the infection at the target site. So far, a simple, fast, precise and  
318 accurate MEPS-UHPLC-PDA method for the determination of MERO,LEVO and LINZ in human  
319 plasma of critical care patients has been developed and applied. Nowadays no paper has been  
320 published for the determination of these drugs. MEPS in combination with UHPLC-PDA has shown  
321 to be adequate for the determination of MERO,LEVO and LINZ in plasma sample achieving good  
322 LOQs and high recovery.

323 About 100 extraction cycles were carried out using the same MEPS sorbent with no loss in  
324 performance. This novel microextraction technique has demonstrated to be simple, cost

325 effectiveness, precise and accurate requiring only small volume of sample and a short run time (5  
326 min).

327

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329

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423 366

## 424 **Figure caption**

425 Figure 1: Chemical structure of Meropenem,(A) Levofloxacin (B) linezolid (C) and the internal  
426 standard ulifloxacin (D, IS)

427 Figure 2: Comparison of extraction recovery percentage obtained from analysis of spiked plasma  
428 samples as a function of the used sorbent (n=5)

429 Figure 3: Mean absolute recovery of MERO, LEVO, LINZ as a function of the extraction cycles and  
430 sample volume.

431 Figure 4: Reproducibility of MERO, LEVO, LINZ and the internal standard (I.S.) as a function of  
432 elution flow rate.

433

434 Figure 5: Chromatograms from human plasma extracts. (a) a blank plasma; (b) a blank plasma  
435 spiked with 2.5 µg/mL of internal standard; (c) a blank plasma spiked at the LOQ value, (d) a blank  
436 plasma spiked with MERO (7.5 µg/mL) LEVO (5 µg/mL) I.S. (2.5 µg/mL) and LINZ (15 µg/mL) and (e)  
437 a human plasma sample collected 4h after an intravenous administration of MERO (Merrem 1g  
438 three times daily) LEVO (Levofloxacin 400mg twice daily) and LINZ (Zyvox 600mg twice daily)  
439 containing MERO (5,7 µg/mL) LEVO (1.2 µg/mL) and LINZ (7 µg/mL). Horizontal axis, retention  
440 time (min).

441

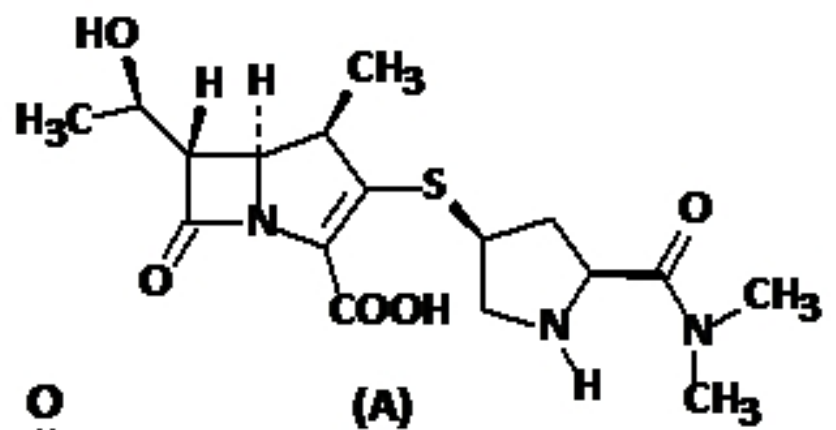
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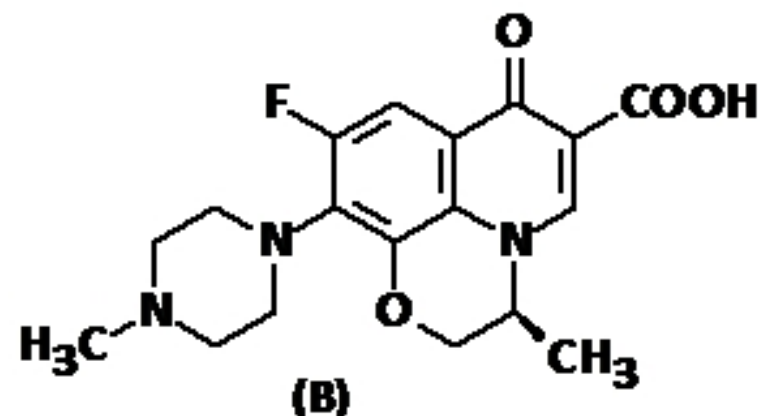
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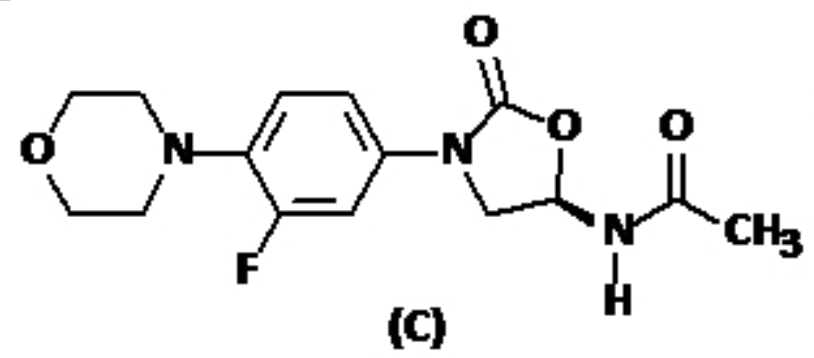
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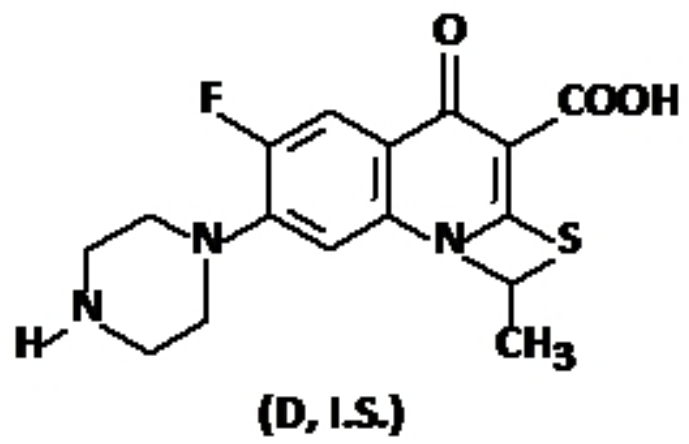
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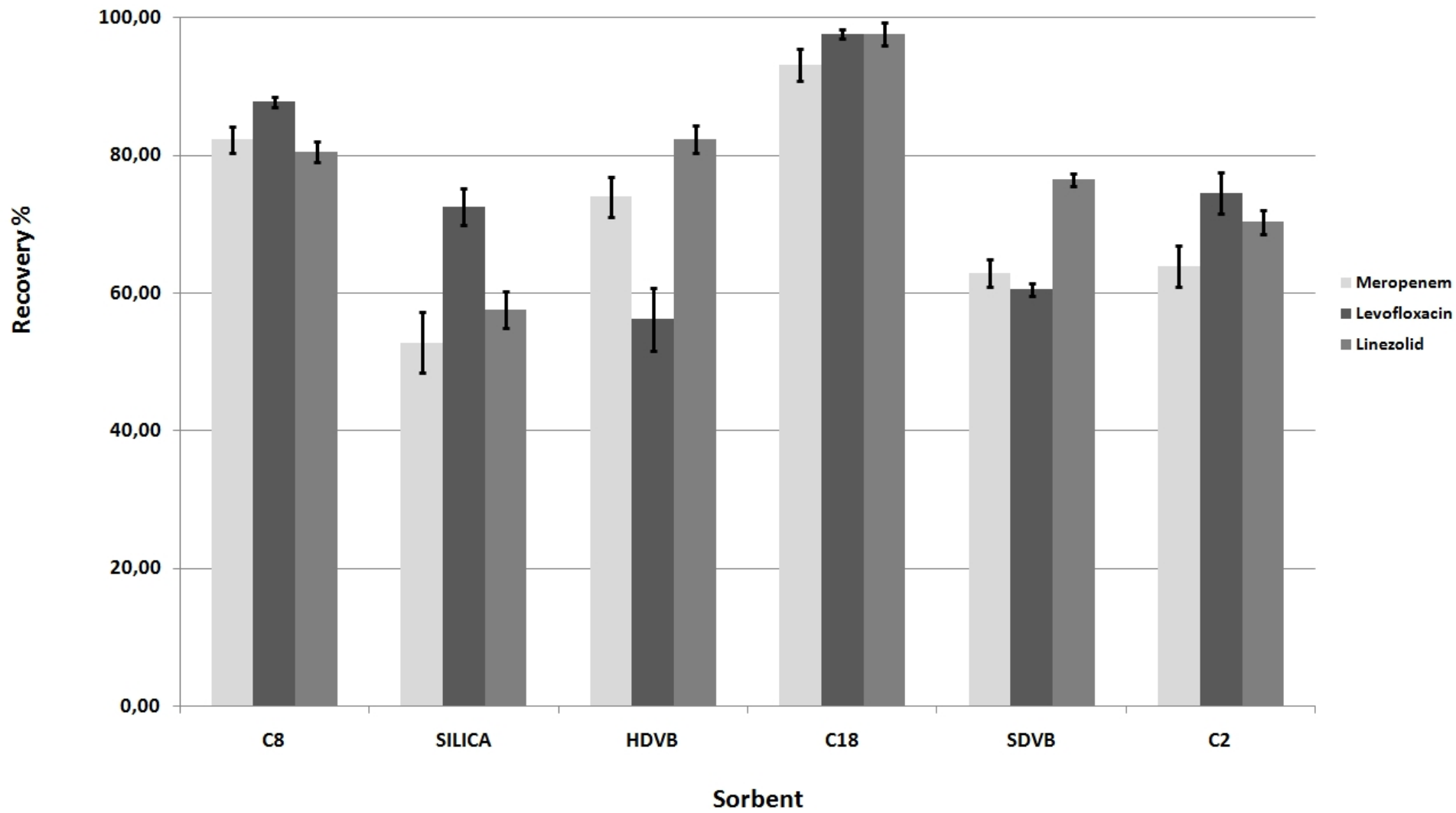
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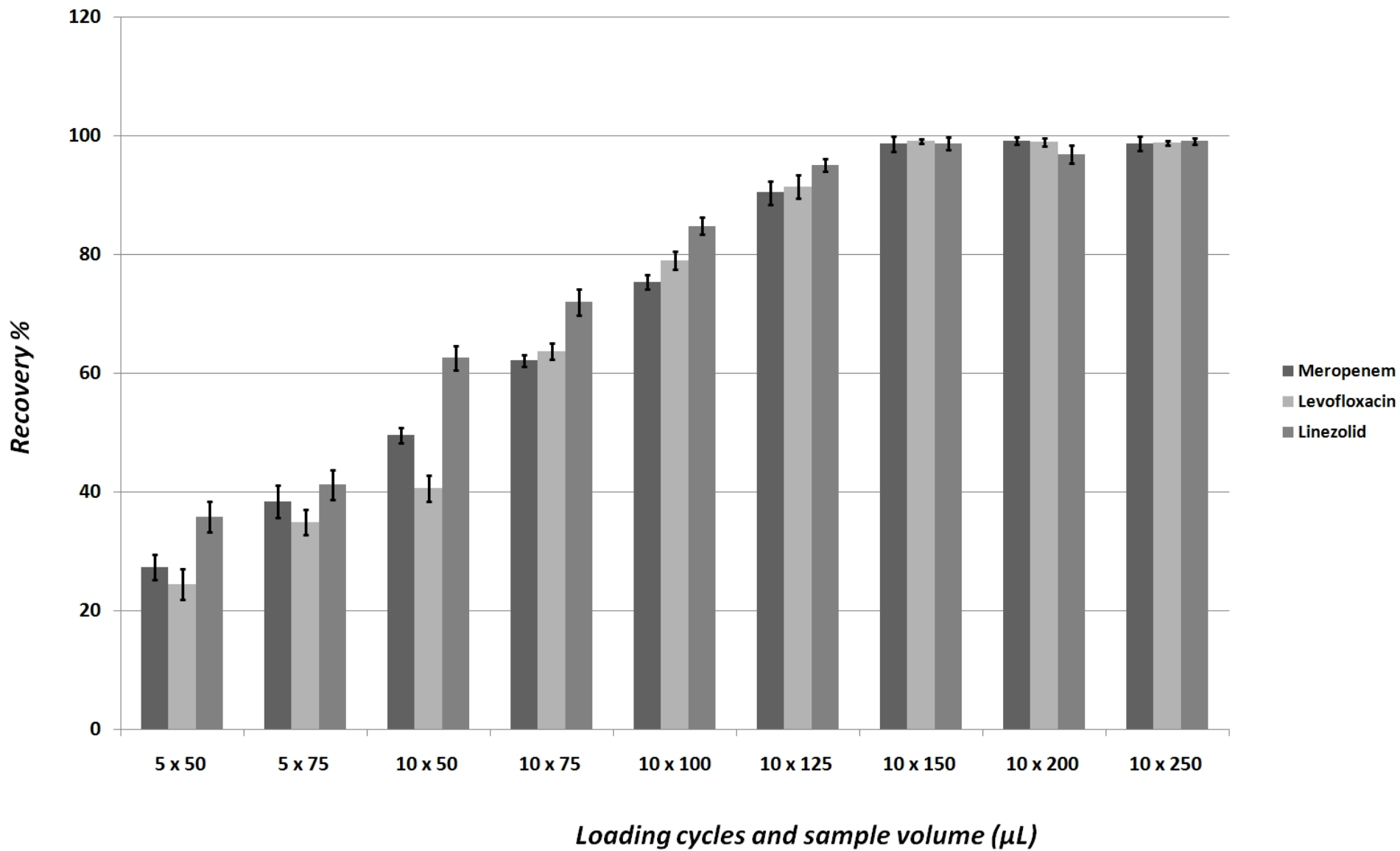


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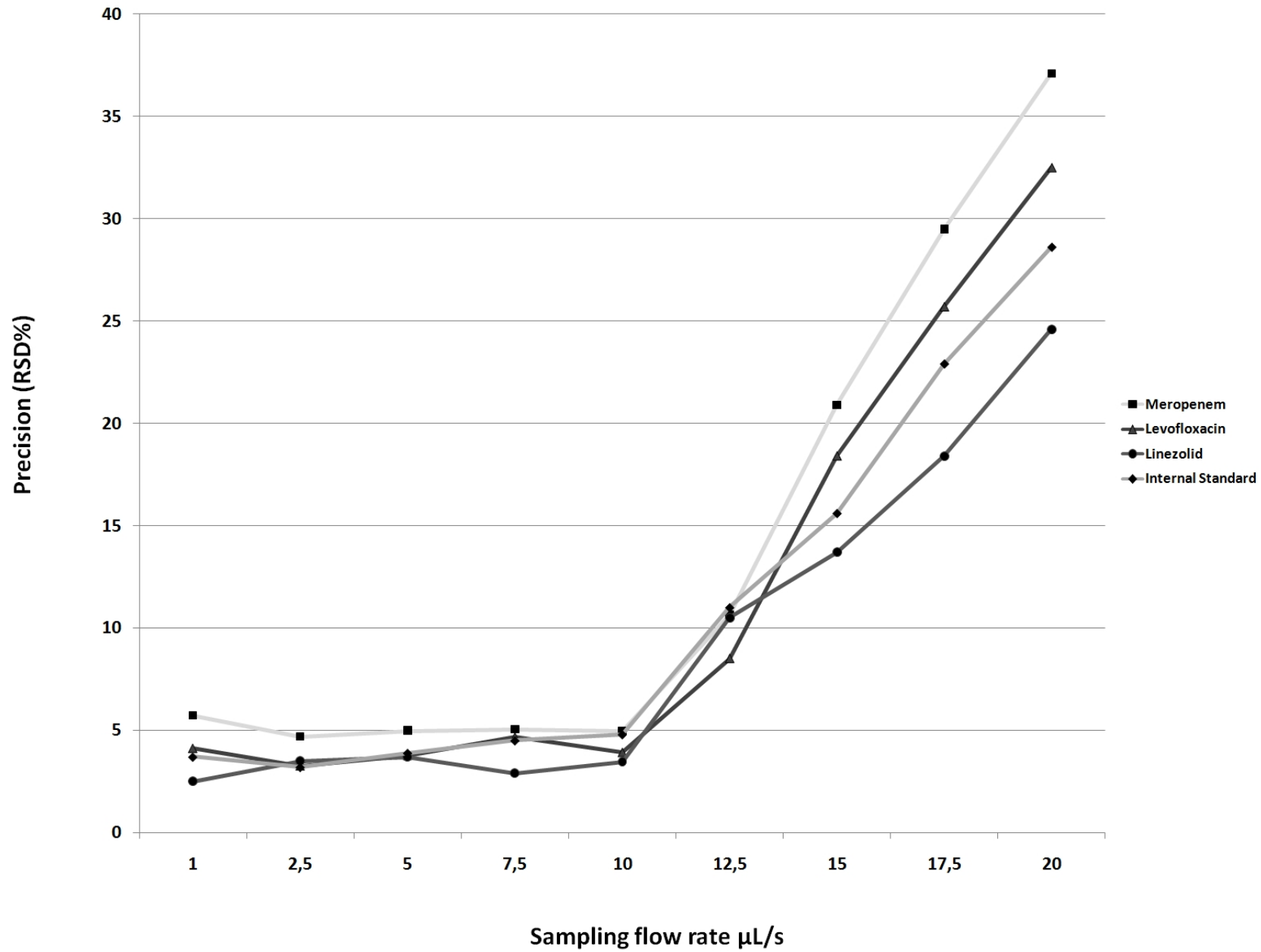


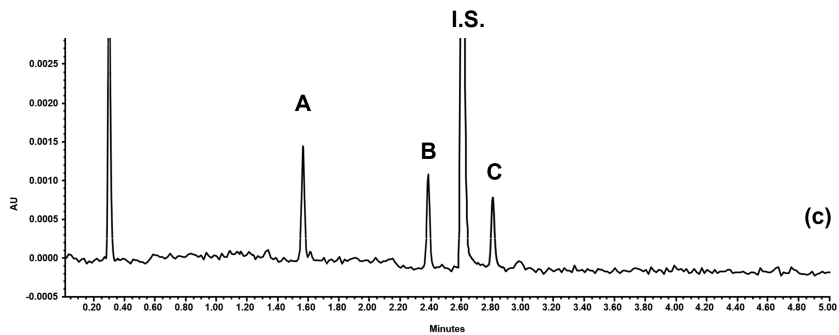
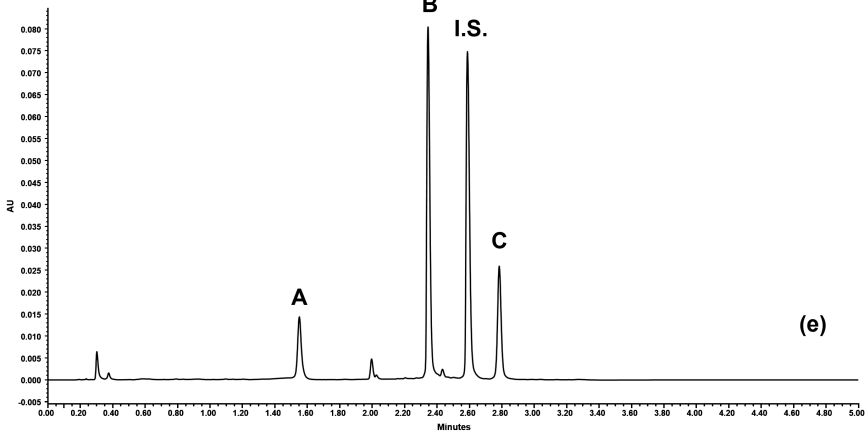
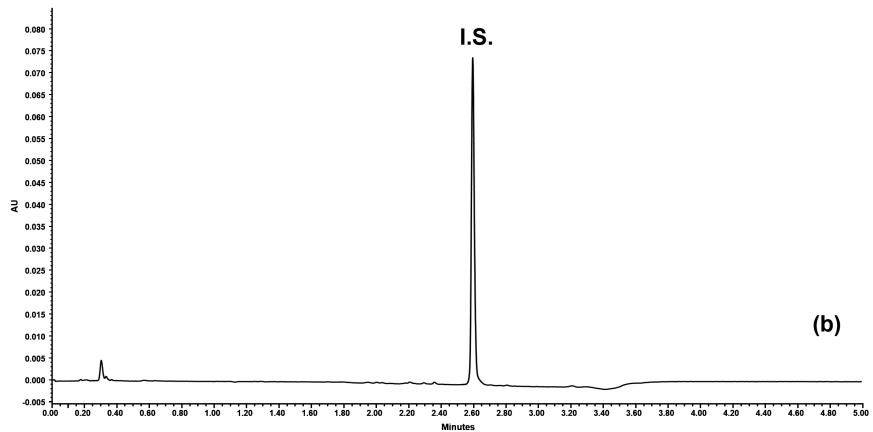
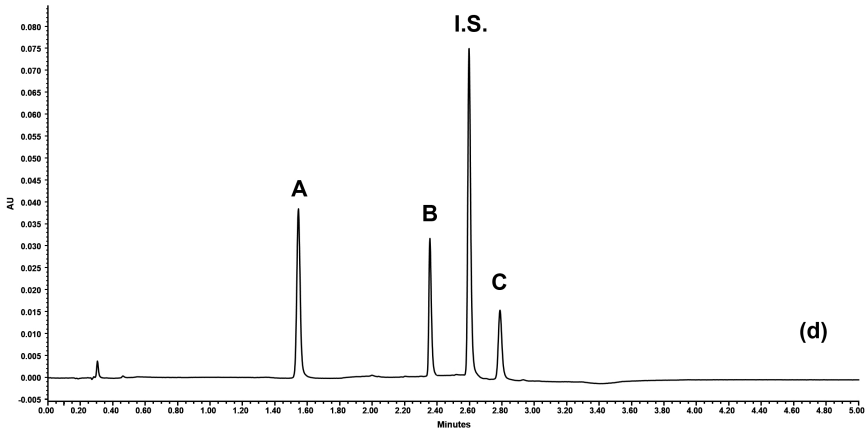
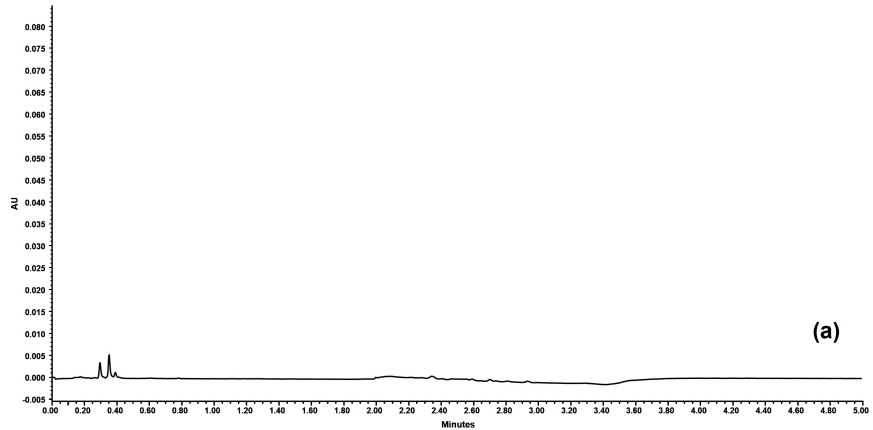
(D, I.S.)











**Table 1:** Summary of the system suitability test for the proposed method

<b>Analyte</b>	<b>RT<sup>a</sup></b>	<b><math>\alpha^b</math></b>	<b>Rs<sup>c</sup></b>	<b>As<sup>d</sup></b>	<b>N<sup>e</sup></b>
<b>Meropenem</b>	1.55	-	12.46	1.11	6051
<b>Levofloxacin</b>	2.36	1.68	3.01	1.08	8911
<b>Ulifloxacin (I.S.)</b>	2.58	1.11	2.84	1.16	13148
<b>Linezolid</b>	2.82	1.14	-	1.09	15708

<sup>a</sup> Retention time (minutes) <sup>b</sup> Selectivity <sup>c</sup> Resolution <sup>d</sup> Asymmetry <sup>e</sup> Theoretical plates

**Table 2:** Intra-day and inter-day accuracy and precision for linezolid and Levofloxacin and meropenem at the LLOQ and QCs value.

	Intra-day (n=6)			Inter-day (n=6)		
	Mean Measured concentration (µg/mL) ± S.D.	Accuracy BIAS (%)	Precision (RSD%)	Mean Measured concentration (µg/mL) ± S.D.	Accuracy BIAS (%)	Precision (RSD%)
<b>Linezolid theoretical concentration (µg/mL)</b>						
0.01 <sup>A</sup>	0.0098 ± 0.0003	-7.4	3.06	0.0099 ± 0.0004	+6.7	4.04
0.02	0.019 ± 0.001	-3.7	5.26	0.018 ± 0.002	-7.8	10.83
1.00	1.03 ± 0.06	+3.4	5.82	1.05 ± 0.06	+5.3	5.71
25.00	24.87 ± 0.57	-0.5	2.29	25.4 ± 1.15	+1.6	4.61
<b>Levofloxacin theoretical concentration (µg/mL)</b>						
0.01 <sup>A</sup>	0.0095 ± 0.002	-4.7	7.42	0.021 ± 0.03	+4.0	7.17
0.02	0.019 ± 0.002	-3.7	10.50	0.021 ± 0.002	+5.9	9.52
0.75	0.74 ± 0.02	-2.2	2.70	0.78 ± 0.07	+4.3	8.97
7.50	7.53 ± 0.19	+0.5	2.52	7.51 ± 0.43	+0.1	5.72
<b>Meropenem theoretical concentration (µg/mL)</b>						
0.02 <sup>A</sup>	0.0191 ± 0.002	-4.3	10.40	0.021 ± 0.03	+4.0	7.17
0.05	0.048 ± 0.003	-3.5	6.25	0.053 ± 0.005	+6.2	9.43
1.00	0.94 ± 0.07	-5.8	7.44	0.78 ± 0.07	+4.3	8.97
25.0	24.1 ± 0.6	+3.6	2.48	26.5 ± 0.9	+6.0	3.39

<sup>A</sup>LLOQ concentration

**Table 3:** Analytical performance of the MEPS-UHPLC-PDA method

	<b>Meropenem</b>	<b>Linezolid</b>	<b>Levofloxacin</b>
$\lambda_{\max}^a$ (nm)	300	254	292
Conc. Range ( $\mu\text{g/mL}$ )	0.02-30	0.01-30	0.01-10
Intercept ( $\pm$ SD)	$2.18 (\pm 0.09) \cdot 10^{-2}$	$7.89 (\pm 0.03) \cdot 10^{-3}$	$4.14 (\pm 0.07) \cdot 10^{-3}$
Slope ( $\pm$ SD)	$0.0182 (\pm 0.008)$	$0.093 (\pm 0.004)$	$0.0182 (\pm 0.008)$
$r^2$	0.9996	0.9998	0.9991
LOQ ( $\mu\text{g/mL}$ )	0.02	0.01	0.01
LOD ( $\mu\text{g/mL}$ )	0.004	0.007	0.004

<sup>a</sup> wavelength of the PDA detection

**Table 4:** Mean recovery of Meropenem, Levofloxacin ,linezolid and the internal standard from human plasma at the QCs concentrations.

<b>Nominal concentration</b>	<b>Mean Measured concentration (<math>\mu\text{g/mL}</math>) <math>\pm</math> S.D.</b>	<b>Recovery (%)</b>	<b>RSD%</b>
<b>MEROPENEM</b>			
0.05	0.0493 $\pm$ 0.003	98.6	6.00
1.00	0.982 $\pm$ 0.08	98.2	8.24
25.00	24.51 $\pm$ 0.34	98.0	1.36
<b>LEVOFLOXACIN</b>			
0.02	0.0191 $\pm$ 0.0012	92.5	6.31
0.75	0.71 $\pm$ 0.05	94.7	7.04
7.50	7.23 $\pm$ 0.44	96.4	5.86
<b>LINEZOLID</b>			
0.02	0.0193 $\pm$ 0.0010	96.5	5.23
1.00	0.96 $\pm$ 0.09	96.7	9.12
25.00	24.10 $\pm$ 0.36	96.4	1.44
<b>INTERNAL STANDARD</b>			
2.50	2.39 $\pm$ 0.05	95.6	2.13

**Table 5:** Plasma concentration of meropenem, levofloxacin and linezolid obtained from real sample subjected to the MEPS-UHPLC-PDA method

Sample	Age	Meropenem ( $\mu\text{g/mL}$ )	Levofloxacin ( $\mu\text{g/mL}$ )	Linezolid ( $\mu\text{g/mL}$ )
#1	72	18.7	1.50	7.8
#2	53	24.1	2.20	22.4
#3	64	9.8	0.98	14.3
#4	67	11.7	3.47	11.6
#5	51	6.1	1.11	9.6

1 **Highlights**

2

3

- A novel sampling and clean-up procedure using microextraction by packed sorbent.

4

- UHPLC-PDA method development and validation

5

- First MEPS-UHPLC-PDA method for determination of meropenem, levofloxacin, linezolid in human plasma

6

7

- Therapeutic drug monitoring in critical care patients