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1	A Validated High Performance Liquid Chromatography – Diode Array Detector
2	Method for the Quantification of Mitomycin C in Plasma, Peritoneal Fluid and Urine
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# 26 Abstract

27 Mitomycin C (MMC) is a quinone-containing alkylating agent, that has been extensively studied in preclinical and clinical work due to its antitumor activity. A thoroughly validated 28 29 high performance liquid chromatography-diode array detector method is provided to quantify 30 MMC in plasma, peritoneal fluid and urine. Porfiromycin served as internal standard. The 31 mobile phase for the plasma, peritoneal fluid and urine analysis consisted of 27% MeOH and 73% 20mM ammoniumacetate buffer (pH 6.5) and 9% ACN and 91% 20mM phosphate buffer 32 33 (pH 6.5) respectively. The residue from 100µL ACN deproteinated plasma was dissolved in 34 250µL mobile phase. Peritoneal fluid and urine were diluted 10-fold in their respective mobile phases. UV detection was performed at 365nm. Quantification of MMC was achieved over a 35 36 linear range of 0.05-5µg/mL and 5-50µg/mL in plasma; 0.1-5µg/mL and 5-100µg/mL in 37 peritoneal fluid; 0.25-5µg/mL and 5-100µg/mL in urine. The limit of quantification was 38 0.05µg/mL in plasma, 0.1µg/mL in peritoneal fluid and 0.25µg/mL in urine. The method was 39 further validated for selectivity, specificity, inter- and intraday precision and accuracy ( $\leq 15\%$ ), 40 extraction recovery and stability. MMC remains stable in the different biofluids for 20 days 41 (short-term stability) at 4°C, -27°C and -80°C and for 80 days (long-term stability) at -80°C. 42

43 Keywords

- 44 Mitomycin C; HPLC-DAD; Plasma; Urine; Peritoneal Fluid
- 45
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- 48

### 49 Introduction

50 Mitomycin C (MMC) is a quinone-containing alkylating agent, isolated from the actinobacteria 51 species, Streptomyces caespitosus [1]. The antibiotic has been extensively studied in preclinical 52 and clinical work due to its antitumor activity. Its most important mechanism of action is 53 through DNA cross-linking. Although MMC is not regarded as a prodrug, it is not active against 54 cancerous tissue as unchanged molecule. The drug is modified as it enters the cell into its active 55 state [2]. It is inactivated by microsomal enzymes in the liver and metabolized in the spleen and 56 kidneys. Today, MMC is widely used for the intraperitoneal (IP) treatment of peritoneal 57 carcinomatosis (PC) from colorectal, appendiceal, ovarian, gastric cancer and diffuse malignant 58 peritoneal mesothelioma [3-8]. Treatment involves a combination therapy of cytoreductive 59 surgery (CRS) and hyperthermic intraperitoneal perioperative chemotherapy (HIPEC). During 60 HIPEC, a heated chemotherapy solution (41-43°C) is instilled in the peritoneal cavity for a 61 predefined time, depending on the chemotherapeutic agent [9]. At present, however, no 62 standardized IP chemotherapy treatment modalities exist, due to variations in choice of drug, 63 dosage, duration, etc. Therefore, there is a pressing need for conducting pharmacologic research 64 towards standardization amongst the myriad of IP treatment protocols currently applied.

65 Throughout literature, various methods have been published regarding the determination and 66 quantification of MMC [10-11]. The most widely used analytical technique to detect MMC in 67 biofluids is reversed phase high performance liquid chromatography (HPLC) combined with 68 UV detection [12-17]. Dalton et al. reported a HPLC method to detect mitomycin C in both 69 human and rat plasma and urine [12]. Using liquid-liquid extraction as sample preparation, a 70 detection limit of 0.5 ng/mL in human plasma could be achieved. Schmid et al. reported the use 71 of a fast and sensitive HPLC-UV method with an online sample clean-up to determine MMC 72 in plasma but data of the development of this analytical method were not published [16]. 73 Eksborg et al. presented a method, using solid phase extraction (SPE) (Sep-Pak C18) as sample

74 preparation, to determine MMC in human plasma and urine within the range of 2-1000 ng/mL 75 and  $0.5 - 4.4 \,\mu\text{g/mL}$ , respectively [17]. All the above-mentioned methods have the limitation of either having time-consuming processing procedures or being reported for only a limited 76 77 number of matrices. Moreover, most publications lack a thorough validation of the described method. Our aim was to develop a validated HPLC-diode array detector (DAD) method to allow 78 79 quantification of MMC in plasma, peritoneal fluid and urine. The described method can easily 80 be used in a routine setting and results are presented of a case study of pharmacologic research 81 regarding the IP administration of MMC in a cancer patient.

#### 82 Materials and methods

#### 83 Safety considerations

When working with chemotherapy, standard safety precautions were applied. These include wearing personal protective equipment (eye protection, protective gloves and protecting clothing) and using standardized handling procedures, including the use of BD PhaSeal<sup>™</sup> closed system transfer devices (Dublin, Ireland), to minimize chemotherapy associated risks. All chemotherapy associated materials and patient samples were disposed in WIVA medical waste containers.

# 90 Chemicals and reagents

MMC, 2 mg, was purchased from Kyowa (Takeda Belgium, Brussels, Belgium) (Figure 1). The
internal standard, porfiromycin, was kindly synthesized by the Laboratory for Organic and Bioorganic Synthesis of Ghent University (Ghent, Belgium) (Figure 1). HPLC-grade acetonitrile
(ACN) and methanol (MeOH) were purchased from VWR International (Fontenay-sous-Bois,
France). Potassium dihydrogen phosphate (KH2PO4) was purchased from Merck Millipore
(Darmstadt, Germany). LC-MS-grade ammonium acetate (C2H7NO2) and syringe filters, 25
mm with 0.45 µm nylon membrane, were purchased from VWR International (Leuven,

Belgium). At all times, ultrapure water of 18.2 MΩcm<sup>-1</sup> was used in the preparations
(Advantage A10 Water Purification System, Merck Millipore, Overijse, Belgium). Nitrogen for
evaporation was purchased from Messer (Messer Group GmbH, Krefeld, Germany). Drug-free
human heparin plasma, urine and peritoneal fluid were kindly provided by the Department of
Laboratory Medicine at the Ziekenhuis Oost-Limburg (Genk, Belgium).

# 103 Calibration standards and validation standards

A standard stock solution of MMC was prepared in ultrapure water at a concentration of 0.4 mg/mL and was stored at room temperature in the dark for up to 1 week. The internal standard solution, porfiromycin, was prepared in MeOH at a concentration of 51.5 mg/mL. Internal standard working solutions were prepared by serially diluting the stock solution in MeOH to reach final concentrations of 10, 100 and 1000  $\mu$ g/mL. The internal standard stock and working solutions were stored in the dark at -80°C and -27°C, respectively.

# 110 Plasma

111 MMC calibration standards in human plasma were prepared by serially diluting the MMC 112 standard stock solution in pooled drug-free plasma to reach final concentrations of 0.05, 0.1, 113 0.5, 1, 5, 10, 20 and 50  $\mu$ g/mL. Quality control samples were prepared by diluting the MMC 114 standard stock solution with pooled drug-free plasma to reach final concentrations of 0.15, 2, 115 5, 16 and 25  $\mu$ g/mL. Calibration standards and quality control samples were freshly prepared 116 each day.

### 117 Peritoneal fluid and Urine

118 Calibration standards in peritoneal fluid and urine were prepared by serially diluting the MMC 119 standard stock solution in drug-free peritoneal fluid and urine to reach final concentrations of 120 0.1, 0.25, 0.5, 1, 5, 10, 50 and 100 µg/mL. Quality control samples were prepared by diluting 121 the MMC standard stock solution with drug-free peritoneal fluid and urine to reach final 122 concentrations of 2, 5, 20 and 80 µg/mL. Calibration standards and quality control samples
123 were freshly prepared each day.

## 124 Sample preparation

# 125 Plasma

Human heparin plasma was obtained by centrifuging blood at 3000 x g for 15 min at 21°C. To 95  $\mu$ L plasma sample volume, 5  $\mu$ L of the internal standard working solution containing 10  $\mu$ g/mL porfiromycin was added. Protein precipitation was performed with 1 mL ACN. After thoroughly mixing on a vortex-mixer for 10 seconds, the mixture was centrifuged at 3000 x g for 10 minutes at 21°C. Subsequently, the supernatant was transferred to a clean glass tube and evaporated to dryness under a steady stream of N<sub>2</sub> at 39°C. The residue was dissolved in 250  $\mu$ L mobile phase and transferred to dark HPLC vials for injection.

# 133 Peritoneal fluid and Urine

Peritoneal fluid and urine were centrifuged at 3000 x g for 10 min at 21°C. To 270  $\mu$ L peritoneal fluid and 95  $\mu$ L urine sample volume, 30  $\mu$ L and 5  $\mu$ L of the internal standard working solutions containing 100  $\mu$ g/mL and 10  $\mu$ g/mL porfiromycin respectively, was added. The matrices were transferred to a clean polypropylene tube, diluted 10-fold in mobile phase, subsequently filtered through 0.45  $\mu$ m nylon syringe filter and transferred to dark HPLC vials before injection.

# 139 Instrumentation and chromatographic conditions

The HPLC system consisted of a Hitachi LaChrom Elite (VWR) system equipped with a
Hitachi L-2200 autosampler, a Hitachi L-2100/2130 pump, a Hitachi L-2300 column oven and
a Hitachi L-2455 DAD (VWR, Leuven, Belgium). The analytical column was a reversed phase
Polaris 3 C18-A, 150 x 3.0 mm, 3 μm particle size (Agilent Technologies, Brussels, Belgium).
The mobile phase for the plasma and peritoneal fluid analysis consisted of 27% MeOH in 73%
20 mM ammoniumacetate buffer (pH 6.5). The mobile phase for the urine analysis consisted

of 9% ACN in 91% 20 mM phosphate buffer (pH 6.5). Both mobile phases were run isocratically at a flow rate of 0.4 mL/min, to ensure back pressures below the threshold of 300 bar. The injection volume was set at 50  $\mu$ L, the plasma and peritoneal fluid analysis were performed at 25°C and the urine analysis at 30°C. The DAD detector was set to measure absorbance at 365 nm. Analytical column care was performed each day by rinsing the column with ultrapure water for 10 column volumes, followed by ACN for approximately 20 column volumes.

# 153 Method validation

154 The described HPLC-DAD method was validated for the following parameters according to the155 generally accepted guidelines for method validation [18].

156 *Linearity* 

157 Linearity was assessed using the previously described calibration standards in plasma, 158 peritoneal fluid and urine. The ratio of the areas of MMC and the internal standard, 159 porfiromycin, was plotted against the analyte concentration and fitted in a linear manner.

160 *Limit of Detection* 

161 Limit of detection (LOD) was defined as the concentration of MMC in the different biofluids162 at which the signal-to-noise ratio corresponded to at least 3.

### 163 Limit of Quantification

Limit of quantification (LOQ) was defined as the concentration of MMC at which the signalto-noise ratio corresponded to at least 10, that could be measured with an acceptable precision and accuracy. Precision was expressed as the relative standard deviation (RSD%); calculated as ((SD/mean) x 100). The accuracy was expressed as the relative error (RE%); calculated as ((found concentration – theoretical concentration)/theoretical concentration) x 100). 169 Acceptable precision and accuracy (n=10) was defined as RSD and RE  $\leq 15\%$  or  $\leq 20\%$  at the 170 lowest level, i.e. the LOQ [19].

# 171 Specificity and Selectivity

The possible interference of endogenous matrix components with the HPLC-DAD assay was evaluated. Resulting chromatograms of plasma, peritoneal fluid and urine samples containing neither MMC nor internal standard were compared with the analysis of the biofluids containing their respective LOQ. Analysis were performed with biofluids from 5 different donors, including two HIPEC treated patients.

#### 177 Precision and Accuracy

178 Interday (n=10) and intraday (n=10) precision and accuracy were evaluated at different 179 validation levels; 0.15, 2, 5, 16 and 25  $\mu$ g/mL MMC in plasma; 2, 5, 20 and 80  $\mu$ g/mL MMC 180 in peritoneal fluid and urine. Acceptable precision and accuracy was defined as RSD and RE 181 <15%.

### 182 Extraction recovery

Extraction recovery of MMC in plasma, peritoneal fluid and urine was evaluated at three different validation levels by comparing the areas of the MMC signal in the respective biofluids with the areas of the MMC signal measured in the corresponding mobile phases. Quality control samples of MMC in matrix were prepared by diluting the MMC standard stock solution in drugfree biofluids to reach final concentrations of 1, 10 and 50  $\mu$ g/mL MMC in plasma; 1, 50 and 100  $\mu$ g/mL MMC in peritoneal fluid and urine. For each concentration, three independent samples were analyzed.

190 Stability

Stability of MMC in the biofluids was assessed at three different validation levels; 2, 16 and 25
µg/mL MMC in plasma; 2, 20 and 80 µg/mL MMC in peritoneal fluid and urine and expressed

in terms of recovery as compared to freshly prepared samples. Short-term and long-term
stability were evaluated by preserving the different samples at 4°C, -27°C and -80°C for 20
days and 80 days respectively. For each concentration, three independent samples were
analyzed.

# 197 Case study

The described HPLC-DAD method was employed in a clinical pharmacokinetic case study. A patient diagnosed with PC from colorectal origin was treated with the CRS and HIPEC procedure. During the 90-minute HIPEC, a total dose of 35 mg/m<sup>2</sup> MMC was administered using the 'Dutch High Dose Mitomycin C Regimen, Triple Dosing Regimen' [20]. At each 15minute time interval, plasma, peritoneal fluid and urine were sampled and processed as previously described. Samples were stored at -80°C until day of analysis.

#### 204 Results

## 205 Chromatography

The total runtime for the plasma (Figure 2a) and peritoneal fluid (Figure 2b) analysis was 12 min. MMC eluted with a retention time of 6.9 min and the internal standard eluted with a retention time of 9.4 min. The total runtime for the urine (Figure 2c) analysis was 30 min, with a retention time of 14.9 min and 23.0 min for MMC and porfiromycin respectively. All peaks exhibit a symmetric shape with only limited tailing. MMC displays two signals at 3.97 min and 6.83 min when measured in the plasma and peritoneal fluid mobile phase (Figure 2d) and at 6.21 min and 15.34 min when measured in the mobile phase for the urine analysis (Figure 2e).

#### 213 Validation

# 214 *Linearity*

- 215 Calibration curves of MMC in plasma, peritoneal fluid and urine were divided into two sections
- 216 to cover a large linear range. The HPLC-DAD method covered a linearity range of 0.05 5
- $217 \mu g/mL$  and  $5 50 \mu g/mL$  in plasma with typical standard curves of y = 2.66x 0.041 with a
- 218 coefficient of determination ( $r^2$ ) of 0.9998 and y = 2.76x 0.818,  $r^2 = 0.9997$  respectively.
- 219 Linearity in peritoneal fluid ranged from  $0.1 5 \mu g/mL$  and  $5 100 \mu g/mL$  with typical standard
- 220 curves of y = 0.11x 0.0017,  $r^2 = 0.9999$  and y = 0.11x 0.0021,  $r^2 = 0.9997$  respectively.
- 221 Linearity in urine ranged from  $0.25 5 \,\mu g/mL$  and  $5 100 \,\mu g/mL$  with typical standard curves
- 222 of y = 0.31x 0.0331,  $r^2 = 0.9995$  and y = 0.34x 0.2699,  $r^2 = 0.996$  respectively.
- 223 Limit of detection
- The LOD was 5 ng/mL MMC in plasma and 20 ng/mL MMC in peritoneal fluid and urine.

#### 225 Limit of Quantification

- 226 The LOQ was 0.05 μg/mL MMC in plasma (n=10; RSD: 5.53%, RE: -15.07%), 0.1 μg/mL
- 227 MMC in peritoneal fluid (n=10; RSD: 2.09%, RE: 8.30%) and 0.25 μg/mL MMC in urine (n =
  228 10; RSD: 6.45%, RE: 14.97%).

#### 229 Specificity and Selectivity

Representative chromatograms of drug-free plasma, peritoneal fluid and urine analysis together with chromatograms of the respective biofluids spiked with their LOQ, evaluating specificity and selectivity are depicted in Figure 2. All samples are from a HIPEC treated patient. We confirm that there was no interference of endogenous matrix components with the HPLC-DAD assay.

### 235 *Precision and Accuracy*

236 The inter- and intraday precision and accuracy were investigated at different MMC levels; 2, 237 16 and 25 µg/mL MMC in plasma; 2, 20 and 80 µg/mL MMC in peritoneal fluid and urine 238 (Table I). Both the RSD and RE are less than 8.94% for the HPLC-DAD analysis of all 239 biofluids. As MMC concentrations in plasma are very low in a cancer patient treated with 240 MMC-based HIPEC (see section case study), an additional level of  $0.15 \,\mu$ g/mL MMC in plasma 241 was evaluated for accuracy and precision. RSD and RE were less than 9.72%. Inter- and 242 intraday precision and accuracy were also validated at a fourth level, 5 µg/mL MMC, i.e. at the 243 intersection of the calibration curves in the respective biofluids, using both calibration curves 244 (Table II). Both the RSD and RE are less than 9.58% for all biofluids.

#### 245 *Extraction recovery*

The extraction recovery from plasma, peritoneal fluid and urine using the sample preparation methods described in the materials and methods section is presented in Table III. In summary, the recovery was assessed at three different MMC levels; 1, 10 and 50  $\mu$ g/mL in plasma; 1, 50 and 100  $\mu$ g/mL in peritoneal fluid and urine. The reported recoveries range from 82.13 – 99.44%.

#### 251 *Stability*

252 The short-term and long-term stability at 4°C, -27°C and -80°C of MMC in plasma, peritoneal 253 fluid and urine is summarized in Table IV. Stability was assessed at three different MMC levels; 254 2, 16 and 25 µg/mL MMC in plasma; 2, 20 and 80 µg/mL MMC in peritoneal fluid and urine, 255 and expressed in terms of recovery. For all biofluids, recovery ranged from 76.61 - 112% after 256 20 days of storage at all defined storage conditions. As to long-term stability; when stored at 257 4°C and -27°C, recoveries ranged from 26.52 – 93.25% with high SDs. When stored at -80°C, recoveries of all biofluids ranged from 76.74 - 103%. Freeze-thaw stability was not investigated 258 259 as part of this validation study. For the application of conducting pharmacologic research regarding the intraperitoneal administration of mitomycin C to cancer patients, the biological samples are only thawed once before analysis and discarded afterwards. Therefore, freeze-thaw stability was not relevant for the particular application.

263 Case study

264 The concentration versus time curve of MMC in plasma, peritoneal fluid and urine is depicted 265 in Figure 3. MMC was quantified in the respective biofluids sampled at each 15-min time 266 interval during the HIPEC procedure. The peritoneal fluid concentration curve demonstrates 267 the administration of MMC in three phases, 'Triple Dosing Regimen', and the subsequent 268 decrease in MMC concentration. The concentration in peritoneal fluid ranged from 6.02 - 9.90269 µg/mL. Plasma concentration, 15 min after initial administration of MMC, was 0.08 µg/mL and 270 increased to 0.31 µg/mL after the 90-minute HIPEC procedure. MMC concentration in urine ranged from 0.25 – 9.95 µg/mL. Representative chromatograms of the MMC and internal 271 272 standard signals, 15 minutes after intraperitoneal chemotherapy administration in the different 273 biofluids of the case study's patient, are presented in Figure 4.

# 274 Discussion

The majority of pharmacokinetic studies reported today draw conclusions from analytical analysis that were developed without a supporting validation. To address this issue, we have demonstrated the use of a thoroughly validated, rapid and sensitive HPLC-DAD method to quantify MMC in plasma, peritoneal fluid and urine.

In literature, several sample preparation methods have been described for MMC quantification in plasma and urine, including liquid-liquid extraction [21], protein precipitation [14] and SPE methods [17]. The evaporation of organic solvents during liquid-liquid extraction methods requires high temperatures and when performed at ambient temperature are often timeconsuming (presumably in the hours range). Moreover, sample preparation methods involving 284 temperatures higher than 40°C will result in the decomposition of MMC and can therefore not 285 be utilized [21]. In the optimization part of this assay, we explored protein precipitation as 286 plasma sample preparation, followed by evaporation at 39°C under a steady stream of N<sub>2</sub>. The 287 improvement we report relative to the protein precipitation method described by Joseph et al. 288 [14] is the time needed for evaporation after precipitation with ACN, i.e. 20 minutes for 36 289 samples as compared to 4-5 hours for 20 samples. For the urine analysis, we explored SPE 290 using different stationary phases; Bond Elut Plexa (30 mg, 3 mL, Agilent Technologies, USA) 291 and ProElut NH<sub>2</sub> (500 mg, 3mL, Dikma Technologies, USA) as sample preparation method. 292 However, SPE sample preparation did not result in an acceptable LOQ and extraction recovery. 293 The above-mentioned sample preparation methods were all optimized aiming at the use of a 294 uniform mobile phase and chromatographic conditions for all biofluids. However, for the urine 295 analysis, the use of 27% MeOH and 73% 20 mM ammoniumacetate buffer (pH 6.5) did not 296 result in acceptable selectivity, specificity and sensitivity. A mobile phase consisting of 9% 297 ACN and 91% 20mM phosphate buffer (pH 6.5) with a runtime of 30 minutes was necessary 298 to achieve good baseline separation for the MMC and internal standard signals in urine and to 299 eliminate interference from contaminants. We can conclude that precipitation with ACN as 300 sample preparation for plasma and a 10-fold dilution in mobile phase followed by filtration 301 through a 0.45 µm syringe filter as sample preparation for peritoneal fluid and urine resulted in 302 limited to no loss of sample and that good baseline separation was achieved used the above-303 mentioned mobile phases.

MMC displays two signals when measured in the mobile phases of the biofluids (Figure 2d and 2e). From liquid chromatography – mass spectrometry (LC-MS) analysis, we know that both signals have the same parent ion (m/z 335) and generate the same fragment ion (m/z 242). Therefore, it can be postulated that two stereoisomers of MMC are present in the reference standard and are retained differently on the chromatographic column. Throughout the validation of the HPLC-DAD method, we performed measurements on the signal with a retention time of
6.83 min (plasma and peritoneal fluid analysis) and 15.34 min (urine analysis); i.e. the
stereoisomeric form of MMC most present in the reference standard.

312 The above described method was validated and proven to be robust in terms of linearity, 313 selectivity, specificity, accuracy, precision and extraction recovery. Regarding stability, 314 recovery of the peritoneal fluid analysis after long-term storage at -80°C was lower than 315 reported for the other biofluids, i.e. 76.74 - 79.35% versus 94.10 - 103%. However, recovery 316 is still in the acceptable range and therefore we confirm that long-term storage of all biofluids, 317 up to 80 days, should be conducted at -80°C. The LOQ we report in plasma is higher than the 318 LOQ reported for the HPLC-UV method by Joseph et al. [14]. However, the presented LOQ's 319 in the different biofluids allow to perform pharmacokinetic studies in which patients receive 320 the 'Dutch High Dose Mitomycin C Regimen, Triple Dosing Regimen', the most widely used 321 HIPEC MMC dosing regimen. The latter is again confirmed by previous pharmacokinetic 322 research reporting on the concentration of MMC administered during HIPEC. Sugarbaker et al. treated patients with low dose MMC, 12.5 mg/m<sup>2</sup> in males and 10 mg/m<sup>2</sup> in females using a 323 324 single administration and diluted this dose in varying volumes of carrier solution. The 325 concentration reported in plasma ranged from 0.01 µg/mL at time point zero to approximately 326 0.5 µg/mL and in peritoneal fluid from 5 µg/mL to approximately 10 µg/mL [22]. Van der 327 Specten *et al.* treated patients with a combination therapy of  $15 \text{ mg/m}^2$  doxorubicin and 15328 mg/m<sup>2</sup> MMC. MMC plasma concentration ranged from 0.1  $\mu$ g/mL at time point zero to 0.25 329  $\mu$ g/mL, MMC urine concentration ranged from 0.1  $\mu$ g/mL to approximately 5  $\mu$ g/mL and from 330  $5 \,\mu\text{g/mL}$  to approximately 10  $\mu\text{g/mL}$  in peritoneal fluid [4].

# 331 Conclusion

We have developed a thoroughly validated HPLC-DAD method to quantify MMC in plasma,peritoneal fluid and urine. As compared to what previously has been reported, our HPLC-DAD

334 method does not require the use of complex chromatographic conditions, expensive and time-335 consuming sample preparation methods. Moreover, good specificity and sensitivity could be 336 achieved resulting in the successful application of our analytical method in a pharmacokinetic 337 case study of a cancer patient, receiving MMC-based HIPEC.

338

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345

# 346 **Conflict of interest statement**

347 The authors declare that they have no conflict of interest.

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#### Tables

#### Table I. Interday and intraday precision (RSD%) and accuracy (RE%) in plasma,

446	peritoneal	fluid	and	urine.
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	n=10		Validation parameter	Theoretical concentration MMC (µg/mL)		
			vandation parameter	2	16	25
			Mean observed concentration ( $\mu g/mL$ ) ± SD	$1.94\pm0.15$	$16.15\pm1.22$	$24.42 \pm 1.72$
		Interday	RSD (%)	7.71	7.56	7.05
	Dlasma		RE (%)	-2.86	0.92	-2.31
	1 1031110		Mean observed concentration ( $\mu g/mL$ ) ± SD	$1.86\pm0.13$	$16.02\pm0.58$	$26.36 \pm 1.16$
		Intraday	RSD (%)	6.92	3.65	4.38
			RE (%)	-7.20	0.10	5.43
				2	20	80
			Mean observed concentration ( $\mu$ g/mL) $\pm$ SD	$1.98\pm0.13$	$19.53 \pm 1.32$	$78.70\pm6.18$
		Interday	RSD (%)	6.40	6.75	7.86
	Peritoneal		RE (%)	-0.93	-2.35	-1.62
	fluid		Mean observed concentration ( $\mu g/mL$ ) $\pm$ SD	$2.00\pm0.13$	$17.70\pm0.55$	$86.20\pm7.70$
		Intraday	RSD (%)	6.70	3.13	8.94
			RE (%)	0.24	-11.48	6.36
			Mean observed concentration ( $\mu g/mL$ ) $\pm$ SD	$1.95\pm0.14$	$20.56 \pm 1.65$	$82.75\pm6.95$
		Interday	RSD (%)	7.03	8.02	8.40
	Urine		RE (%)	-2.47	2.80	3.43
	onne		Mean observed concentration ( $\mu g/mL$ ) ± SD	$1.94\pm0.10$	$20.41\pm0.84$	$86.25 \pm 1.41$
		Intraday	RSD (%)	5.01	4.10	1.64
			RE (%)	-2.97	2.04	7.82
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# 456 Table II. Interday and intraday precision (RSD%) and accuracy (RE%) of 5 μg/mL MMC

		Validation parameter	Theoretical concentration MMC (µg/mL) 5	
				10n curves
		Maan absorbed concentration $(\mu g/mL) + SD$	0.03 - 3	5 - 30
	Interday	Mean observed concentration ( $\mu$ g/mL) $\pm$ SD	$3.03 \pm 0.10$	$5.04 \pm 0.2$
	Interday	R5D (70)	0.92	9.58
Plasma		Mean observed concentration $(\mu g/mL) + SD$	$5.18 \pm 0.18$	5.01
	Intradav	RSD (%)	3.44	4.26
	5	RE (%)	3.69	2.16
			0.1 - 5	5 - 100
		Mean observed concentration ( $\mu g/mL$ ) $\pm$ SD	$5.00 \pm 0.02$	4.95 ± 0.2
	Interday	RSD (%)	0.45	5.17
Peritoneal		RE (%)	0.01	-1.04
fluid		Mean observed concentration ( $\mu g/mL$ ) ± SD	$5.09\pm0.05$	$5.05 \pm 0.0$
	Intraday	RSD (%)	0.96	1.22
		RE (%)	1.71	0.94
			0.25 - 5	5 - 100
		Mean observed concentration ( $\mu g/mL$ ) ± SD	$5.01 \pm 0.01$	5.13 ± 0.3
	Interday	RSD (%)	0.12	6.67
Urine		RE (%)	0.13	2.65
UTILIE		Mean observed concentration (µg/mL) $\pm$ SD	$4.85\pm0.07$	$5.14 \pm 0.0$
	Intraday	RSD (%)	1.42	1.27
		RE (%)	-2.98	2.82

# 457 assessed in the different calibration curves in plasma, peritoneal fluid and urine.

	Theoretical concentration MMC (µg/mL)	Extraction recovery (%) ± SD	
	1	$82.32\pm6.16$	
Plasma	10	$87.57\pm2.50$	
	50	$88.42\pm6.64$	
Domiton col	1	$99.44 \pm 3.13$	
fluid	50	$96.55\pm5.06$	
	100	$96.90 \pm 3.57$	
	1	$86.16\pm0.78$	
Urine	50	$93.20 \pm 0.77$	
	100	$86.24 \pm 0.83$	

# 467 Table III. Extraction recovery from plasma, peritoneal fluid and urine.<sup>a</sup>

489	Table IV. Short-term and long-term stability of MMC analysis in plasma, peritoneal fluid
490	and urine at 4°C, -27°C and -80°C. <sup>a</sup>

		Temperature (°C)	Recovery (%) $\pm$ SD ( $n = 3$ ) Theoretical concentration MMC (µg/mL)		
			2	16	25
Plasma	Short-term (20 days)	4	$96.96 \pm 2.85$	$93.72\pm0.14$	$94.47 \pm 2.83$
		-27	$103 \pm 6$	$101 \pm 2$	$92.91 \pm 1.34$
		-80	$112 \pm 6$	$103\pm0.19$	$91.30\pm2.77$
	Long-term (80 days)	4	$69.46\pm6.40$	$47.37 \pm 8.96$	$89.18 \pm 1.28$
		-27	$93.25\pm4.54$	$77.02\pm28.28$	$91.64 \pm 5.34$
		-80	$96.50\pm3.55$	$103 \pm 10$	$98.48 \pm 1.11$
			2	20	80
Peritoneal fluid	Short-term (20 days) Long-term (80 days)	4	$96.96 \pm 2.85$	$93.72\pm0.14$	$81.69\pm0.75$
		-27	$103 \pm 6$	$101 \pm 2$	$95.54\pm0.28$
		-80	$112 \pm 6$	$103\pm0.2$	$99.41 \pm 1.07$
		4	$64.83\pm9.25$	$87.01 \pm 19.23$	$78.11 \pm 9.34$
		-27	$65.37 \pm 3.15$	$73.14\pm0.88$	$77.09 \pm 1.04$
		-80	$78.35\pm0.87$	$76.74 \pm 1.60$	$79.35\pm0.57$
Urine	Short-term (20 days)	4	$76.61 \pm 1.15$	$89.83 \pm 1.06$	$91.17\pm0.81$
		-27	$95.67 \pm 5.29$	$100 \pm 1$	$96.68 \pm 0.63$
		-80	$110 \pm 1$	$109 \pm 1$	$99.32\pm~0.42$
	Long-term (80 days)	4	$27.96\pm0.49$	$38.57 \pm 1.39$	$53.93 \pm 3.15$
		-27	$26.52\pm9.21$	$60.96 \pm 2.84$	$40.97\pm32.82$
		-80	$94.10 \pm 1.27$	$96.90\pm0.82$	$99.35\pm0.86$

<sup>491 &</sup>lt;sup>a</sup> For each concentration, three independent samples were analyzed (n=3).

501 Figure legends

502

- 503 Figure 1. Chemical structure of mitomycin C and porfiromycin.
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505 Figure 2. Selectivity and specificity of the HPLC-DAD mitomycin C analysis in plasma, 506 peritoneal fluid and urine (a) HPLC-DAD analysis of MMC (6.84 min) and the internal 507 standard, porfiromycin (9.28 min), in plasma. Plasma spiked with the LOQ, 0.05 µg/mL MMC 508 and 0.5 µg/mL porfiromycin (blue). Drug-free plasma (green). (b) HPLC-DAD analysis of 509 MMC (7.01) and the internal standard, porfiromycin (9.50 min), in peritoneal fluid. Peritoneal 510 fluid spiked with the LOQ, 0.1 µg/mL MMC and 10 µg/mL porfiromycin (blue). Drug-free 511 peritoneal fluid (green). (c) HPLC-DAD analysis of MMC (14.85 min) and the internal 512 standard, porfiromycin (22.98 min), in urine. Urine spiked with the LOQ, 0.25 µg/mL MMC 513 and 5 µg/mL porfiromycin (blue). Drug-free urine (green). (d) HPLC-DAD analysis of MMC 514 and the internal standard, porfiromycin, in 27% MeOH / 73% 20 mM ammoniumacetate buffer 515 (pH 6.5). The mobile phase of the plasma and peritoneal fluid analysis was spiked with  $5 \mu g/mL$ 516 MMC and 1 µg/mL porfiromycin (9.28 min). MMC displays two signals at 3.97 min and 6.83 517 min. (e) HPLC-DAD analysis of MMC and the internal standard, porfiromycin, in 9% ACN / 518 91% 20 mM phosphate buffer (pH 6.5). The mobile phase of the urine analysis was spiked with 519 5 µg/mL MMC and 1 µg/mL porfiromycin (23.75 min). MMC displays two signals at 6.21 min 520 and 15.34 min.

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522 Figure 3. Concentration versus time graph of intraperitoneal mitomycin C, after 523 administering 'Triple Dosing Regimen' during a 90-minute hyperthermic intraperitoneal 524 perioperative chemotherapy procedure At each 15-minute time interval, plasma (red), 525 peritoneal fluid (blue) and urine (green) were sampled. Concentration of MMC in peritoneal

526	fluid ranged from 6.02 – 9.90 $\mu$ g/mL. Concentration of MMC in plasma ranged from 0.08 -
527	0.31 $\mu$ g/mL. Concentration of MMC urine ranged from 0.25 – 9.95 $\mu$ g/mL.

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Figure 4. Representative chromatograms of the mitomycin C and internal standard signals, 15 minutes after administering 'Triple Dosing Regimen' hyperthermic intraperitoneal perioperative chemotherapy (a) HPLC-DAD analysis of MMC (6.73 min) and the internal standard, porfiromycin (9.11 min), in plasma. (b) HPLC-DAD analysis of MMC (6.77) and the internal standard, porfiromycin (9.15 min), in peritoneal fluid. (c) HPLC-DAD analysis of MMC (12.39 min) and the internal standard, porfiromycin (19.50 min), in urine.