Analytical and Bioanalytical Chemistry

Electronic Supplementary Material

Critical assessment of different methods for quantitative measurement of metallodrug-protein associations

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ICP-QMS iCAP Thermo Scientific operation parameters				
Nebulizer	PFA-ST			
Spray chamber	Cyclonic			
Nebulizer gas flow	1.01 L/min			
Aux. gas	0.99 L/min			
Plasma gas	14 L/min			
Reaction gas	0.370 mL/min			
ICP RF Power	1550 W			
m/z measured	194.97, 47.97			
Thermo Scientific Transcend system chromatographic conditions				
Thermo Scientific Transcend system chromatographic con	nditions			
Thermo Scientific Transcend system chromatographic con HPLC Column	nditions Acquity UPLC Protein BEH SEC, 4.6 x 150 mm, 200A, 1.7 µm, 10kDa- 500kDa, Waters			
	Acquity UPLC Protein BEH SEC, 4.6 x 150 mm, 200A, 1.7 μm, 10kDa-			
HPLC Column	Acquity UPLC Protein BEH SEC, 4.6 x 150 mm, 200A, 1.7 μm, 10kDa- 500kDa, Waters			
HPLC Column Eluent	Acquity UPLC Protein BEH SEC, 4.6 x 150 mm, 200A, 1.7 μm, 10kDa- 500kDa, Waters 50 mM CH ₃ COONH ₄ , pH = 6.0			
HPLC Column Eluent Flow Rate	Acquity UPLC Protein BEH SEC, 4.6 x 150 mm, 200A, 1.7 μm, 10kDa- 500kDa, Waters 50 mM CH ₃ COONH ₄ , pH = 6.0 400 μL/min			

Table S1 ICP-MS operation parameters and chromatographic conditions for the kinetic studies of

 KP2156 and KP2157 in FCS

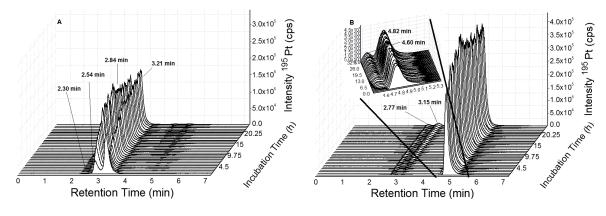


Fig. S1 Chromatographic separations with the column Acquity UPLC Protein BEH SEC, 4.6 x 150 mm, 200A, 1.7 μ m, 10kDa-500kDa, of the platinum complexes (~1 μ M) upon incubation in FCS during 24h at 37°C (corresponding approximately to a molar ratio of drug:albumin of ~1:600). (A) KP2156 shows a high binding to albumin (3.21, 2.84, 2.54 and 2.30 min). The free drug could not be observed. (B) KP2157 shows a lower binding to albumin (3.15 and 2.77 min). The free drug could be observed at 4.82 min. A hydrolysis product was observed in kinetics studies of the drug in water as well as in the incubation in FCS (4.60 min)

Table S2 Analytical figures of merit of the SEC and UHPLC SEC columns

Column	Compound	Retention Time (min)	$N (m^{-1})^{a}$	Η (μm) ^b
BioBasic SEC-60 A, 4.6x250 mm, 5µm, Thermo.	Methionine	12.9	17072	59
Acquity UPLC Protein BEH SEC, 4.6 x 150 mm, 125A, 1.7 μm, Waters.	Methionine	5.34	61960	16

^a Theoretical Plate Number m⁻¹.

^b Theoretical Plate Height µm.

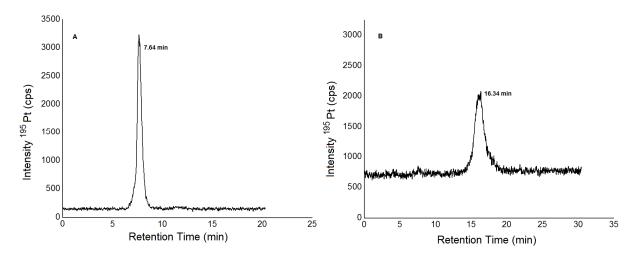


Fig. S2 SEC chromatograms (BioBasic SEC-60 A, 4.6x250 mm, 5 μ m) of (A) an incubation of KP2156 0.1 μ M (20 μ g/L of Pt) 1:10 in FCS during 45 min at 37°C and diluted 1:10 in water prior to the analysis. A high binding of this compound to albumin could be observed (7.64 min). (B) An incubation of KP2157 0.1 μ M (20 μ g/L of Pt) 1:10 in FCS during 45 min at 37°C and diluted 1:10 in water prior to the analysis. This compound showed no binding to albumin appearing mainly as free drug (16.34 min)

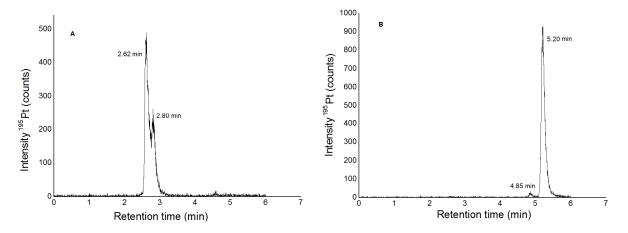


Fig. S3 UHPLC SEC chromatograms (Acquity UPLC Protein BEH SEC, 4.6 x 150 mm, 125A, 1.7 μ m) of (A) an incubation of KP2156 0.1 μ M (20 μ g/L of Pt) 1:10 in FCS during 45 min at 37°C. A high binding of this compound to the monomer and the dimer of albumin could be observed (2.62 and 2.80 min respectively) (B) an incubation of KP2157 0.1 μ M (20 μ g/L of Pt) 1:10 in FCS during 45 min at 37°C. This compound showed no binding to albumin appearing mainly as free drug (5.20 min). A hydrolysis product was observed in the standard of the drug in water as well as in the incubation in FCS (4.85 min)

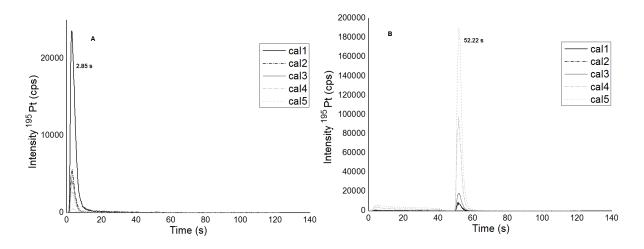


Fig. S4 TFC chromatograms (Fluoro XL 0.5x50 mm) of (A) standards of KP2156 in FCS 1:10 at different concentrations. (B) standards of KP2157 in water at different concentrations