

ELECTRONIC SUPPLEMENTARY INFORMATION

Esterification of free carboxylic group from the lutidinic acid ligand as a tool to improve the cytotoxicity of Ru(II) complexes

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Table S1 Crystal data and structure refinement parameters obtained for the complexes 1 and 2.

	1	2
Empirical formula	[RuC ₄₅ H ₄₀ N ₃ O ₄ P ₂] ₂ PF ₆	[RuC ₄₇ H ₄₄ N ₃ O ₄ P ₂] ₂ PF ₆
Formula weight	994.81	1022.83
Temperature (K)	293(2)	298(2)
Wavelength (Å)	0.71073	0.71073
Crystal system	Monoclinic	Monoclinic
Space group	C2/c	P2 ₁ /n
a (Å)	40.71(2)	13.5183(4)
b (Å)	15.001(9)	12.8666(3)
c (Å)	14.658(9)	28.1285(7)
α (°)	90	90
β (°)	95.496(7)	100.746(10)
γ (°)	90	90
Volume (Å ³)	8910(9)	4806.7(2)
Z	8	4
Density (calculated) (Mg/m ³)	1.415	1.413
Absorption coefficient (mm ⁻¹)	0.523	0.494
F(000)	3864	2088.0
Crystal size (mm ³)	0.26 x 0.26 x 0.20	0.21 x 0.10 x 0.09
Theta range for data collection (°)	1.977 to 27.485	2.719 to 30.508
Index ranges	-52 ≤ h ≤ 52, -19 ≤ k ≤ 19, -18 ≤ l ≤ 18	-19 ≤ h ≤ 19, -18 ≤ k ≤ 18, -40 ≤ l ≤ 40
Reflections collected	39930	73392
Independent reflections	10192 [R(int) = 0.0614]	14670 [R(int) = 0.0394]
Completeness to theta (%)	100.0	100.0
Refinement method	Full-matrix least-squares on F ²	Full-matrix least-squares on F ²
Data / restraints / parameters	10192 / 0 / 563	14670 / 0 / 574
Goodness-of-fit on F ²	1.140	1.098
Final R indices [I > 2σ(I)]	R1 = 0.0563, wR2 = 0.1403	R1 = 0.0548, wR2 = 0.1384
R indices (all data)	R1 = 0.0744, wR2 = 0.1610	R1 = 0.0782, wR2 = 0.1616
Extinction coefficient	0.00268(15)	0.00068(13)
Largest diff. peak and hole (e.Å ⁻³)	0.848 and -0.938	1.36 and -0.95

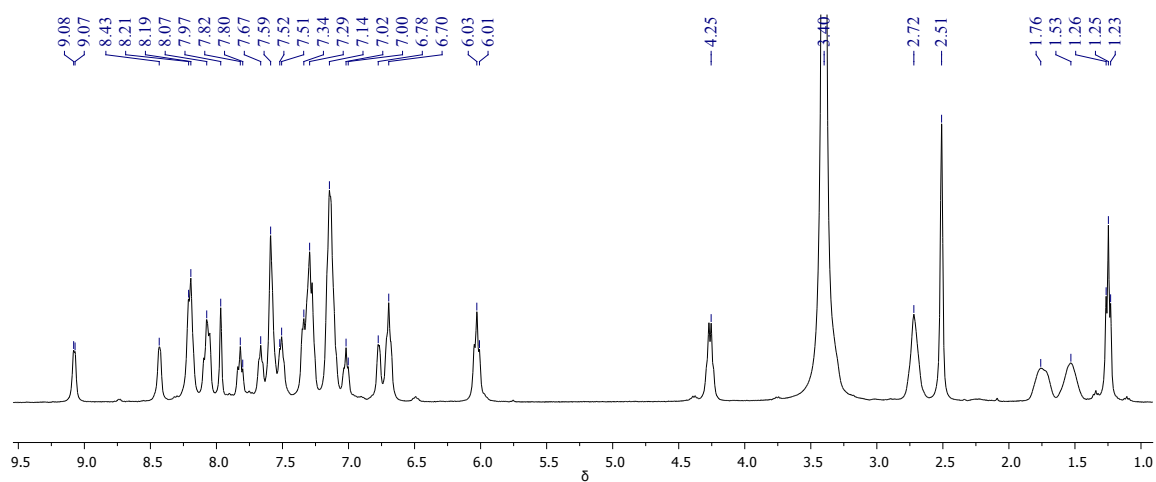


Figure S1 ^1H NMR spectra of the complex 2 with peak assignment (DMSO-d_6)

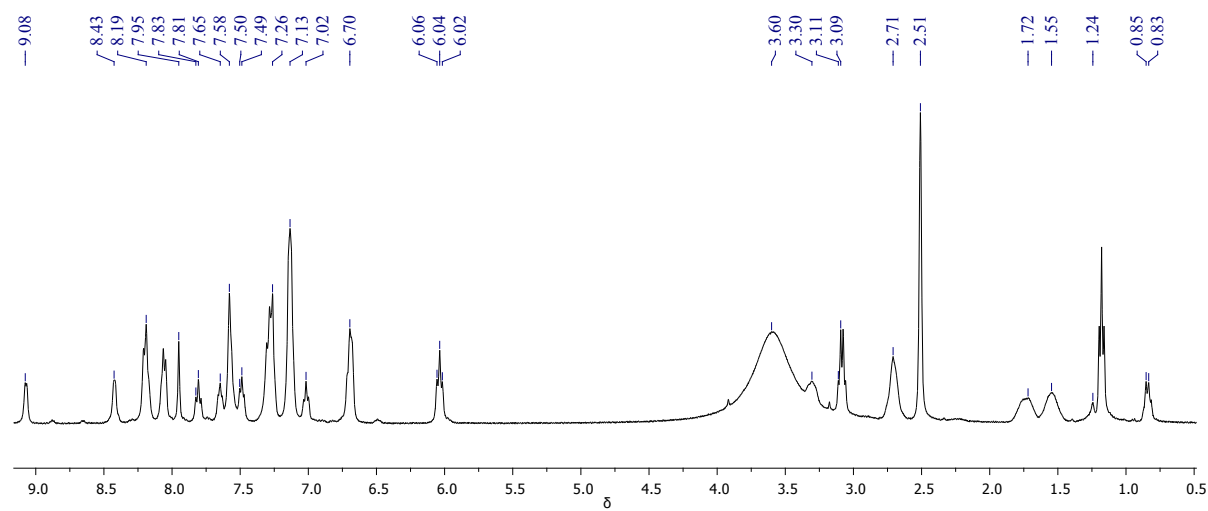


Figure S2 ^1H NMR spectra of the complex 1 with peak assignment (DMSO-d_6)

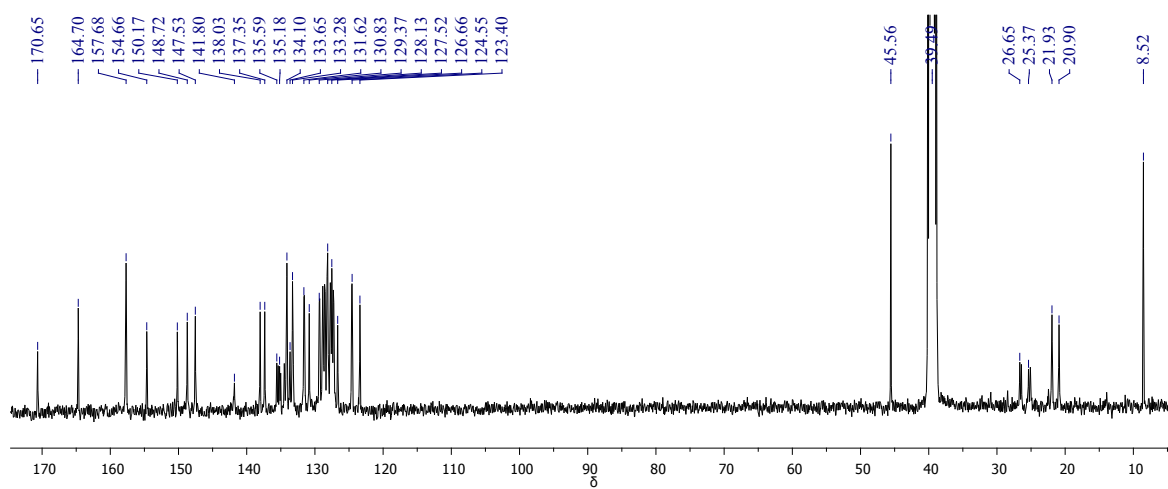


Figure S3 $^{13}\text{C}\{^1\text{H}\}$ NMR spectra of the complex 1 with peak assignment (DMSO-d_6)

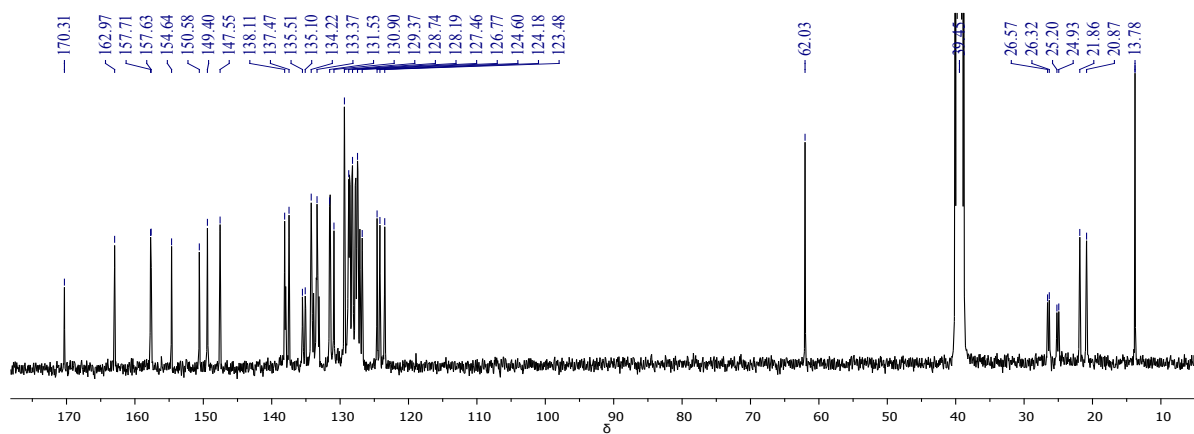


Figure S4 $^{13}\text{C}\{^1\text{H}\}$ NMR spectra of the complex 2 with peak assignment (DMSO-d_6)

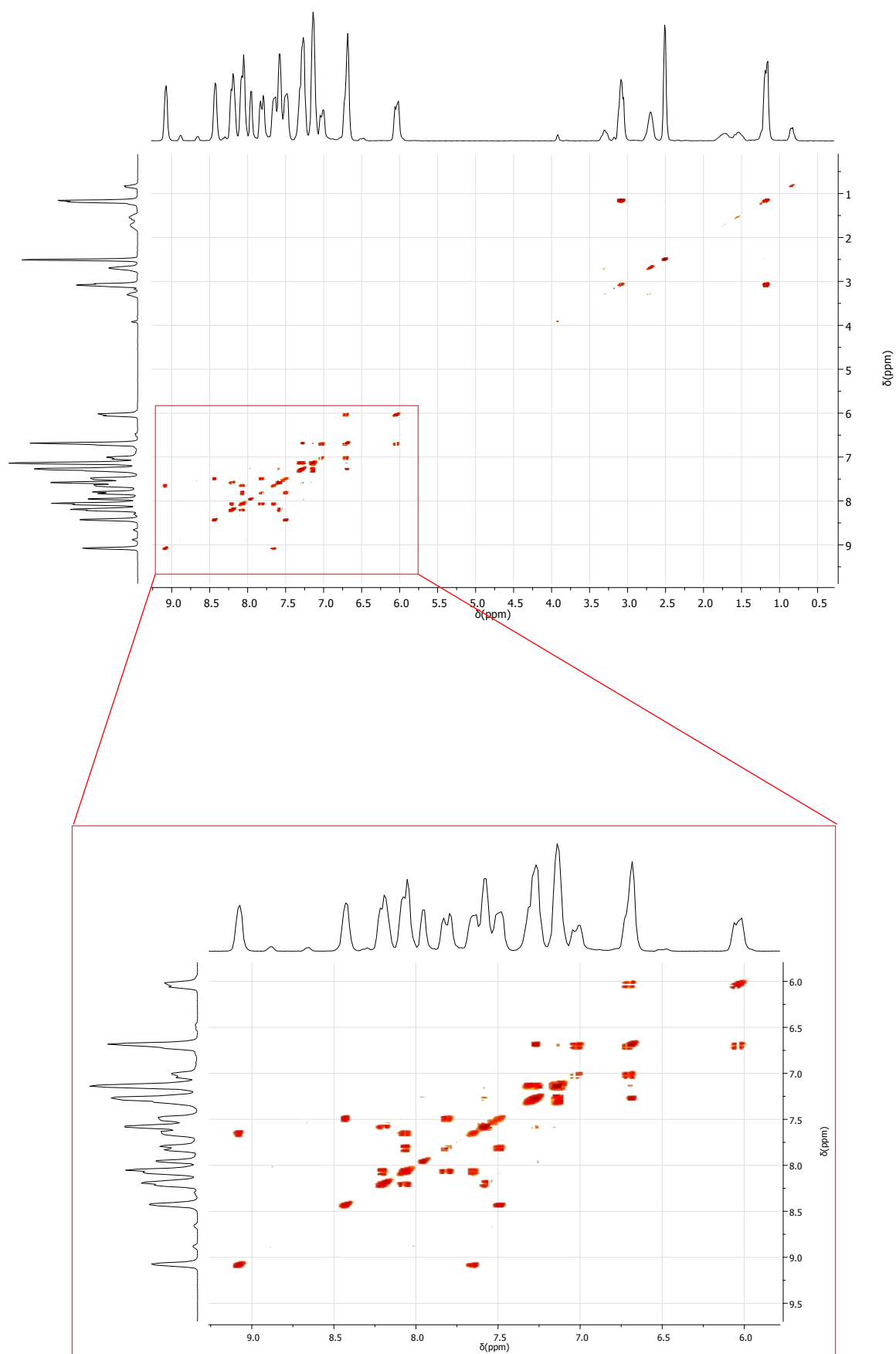


Figure S5 ^1H - ^1H COSY NMR spectra of the complex 1 (DMSO-d_6)

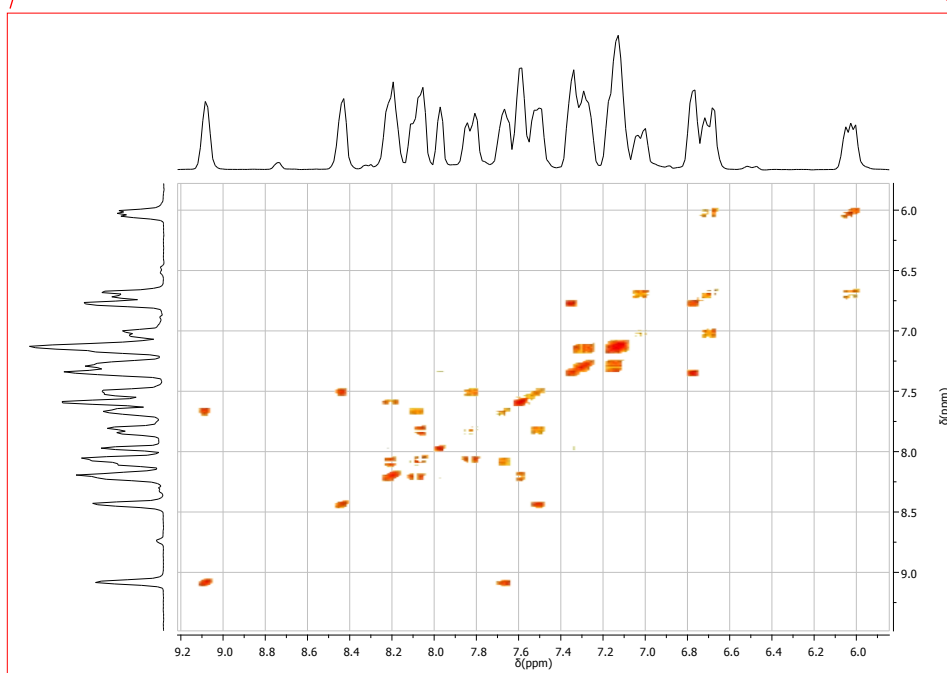
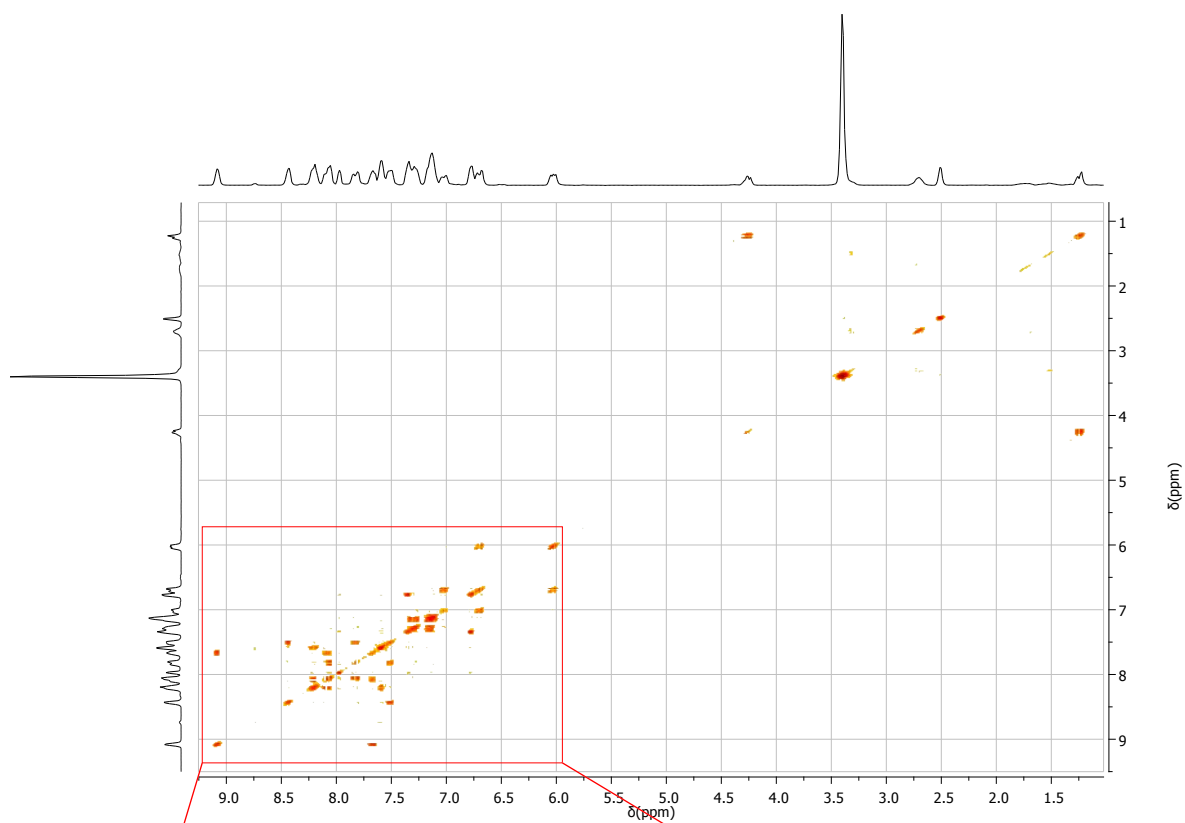


Figure S6 ^1H - ^1H COSY NMR spectra of the complex 2 (DMSO- d_6)

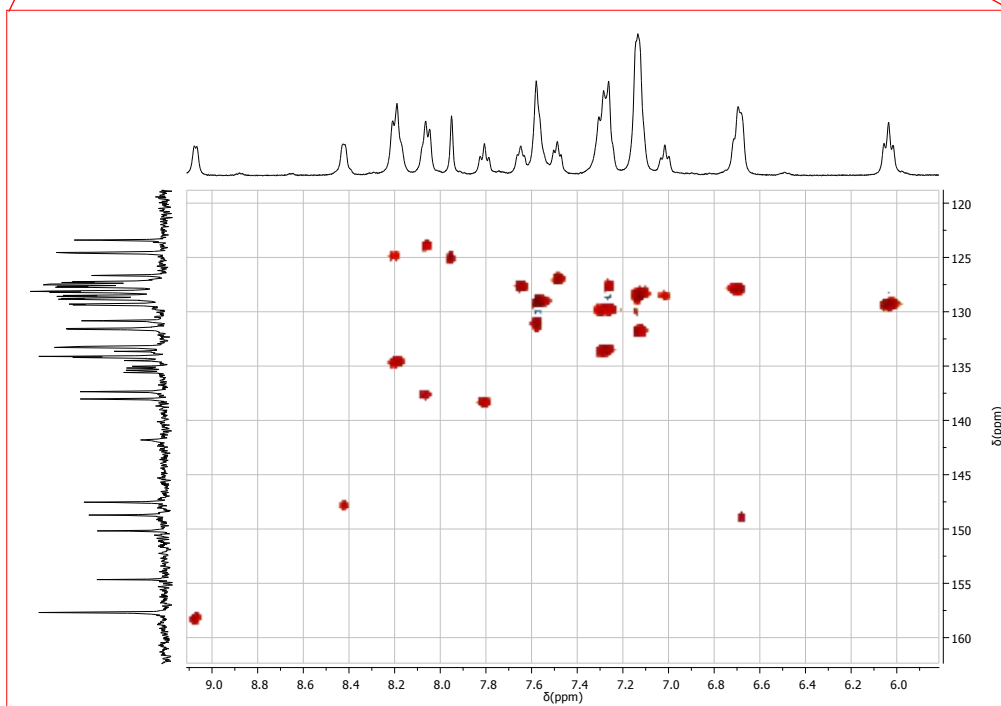
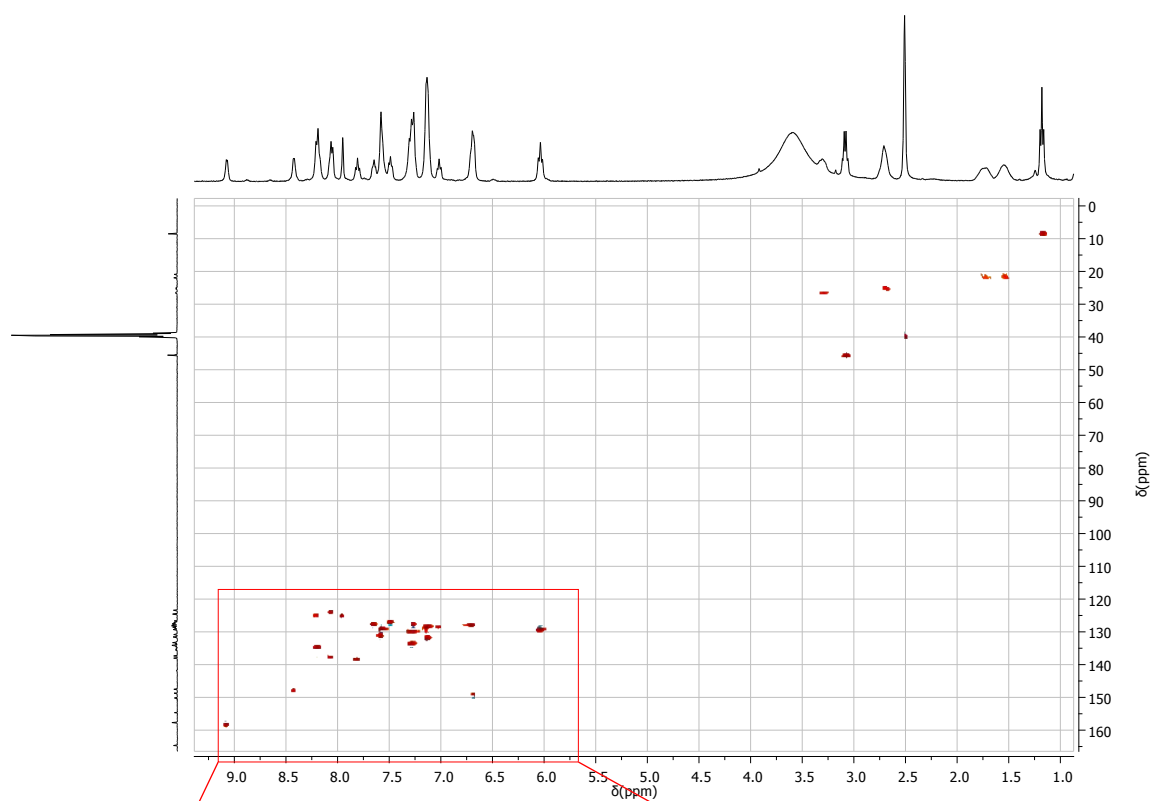


Figure S7 ^1H - ^{13}C HSQC NMR spectra of the complex 1 (DMSO-d_6)

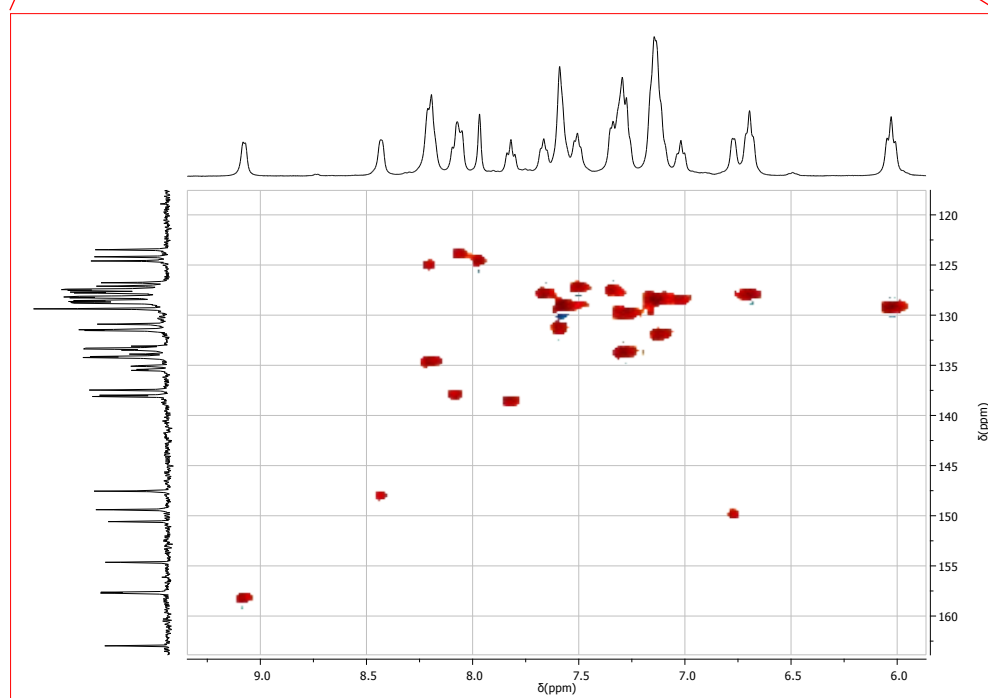
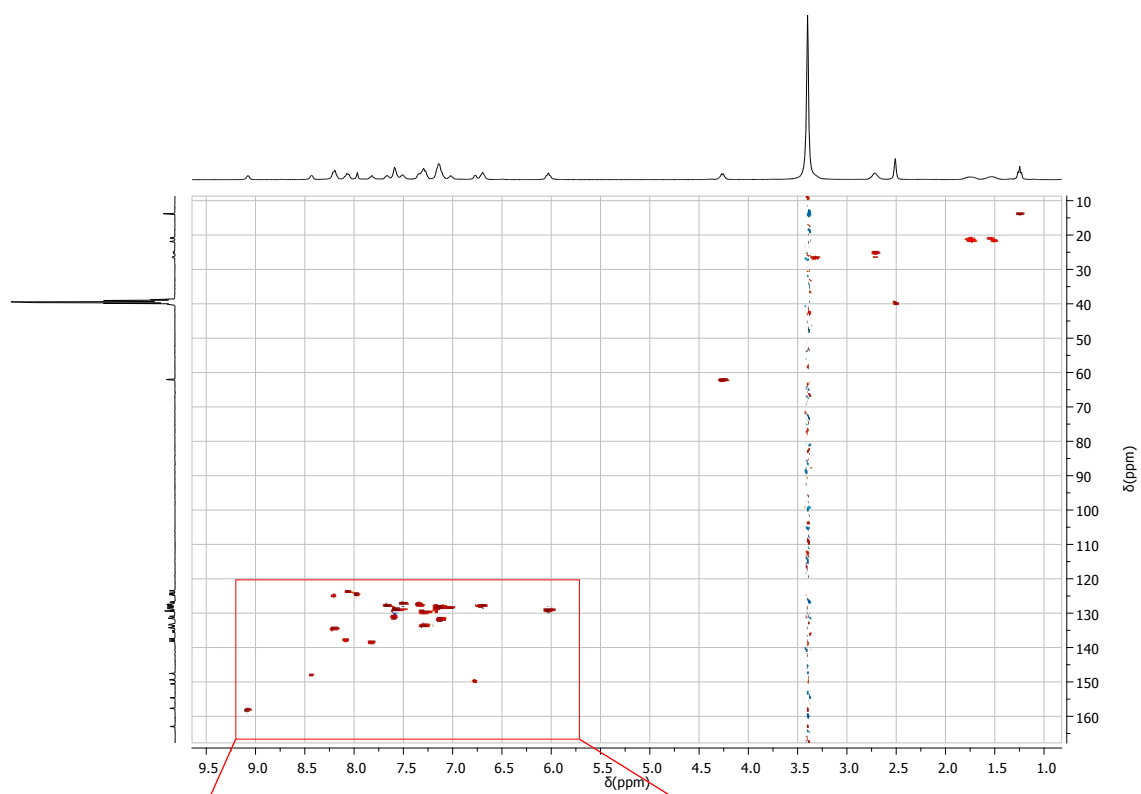


Figure S8 ^1H - ^{13}C HSQC NMR spectra of the complex 2 (DMSO- d_6)

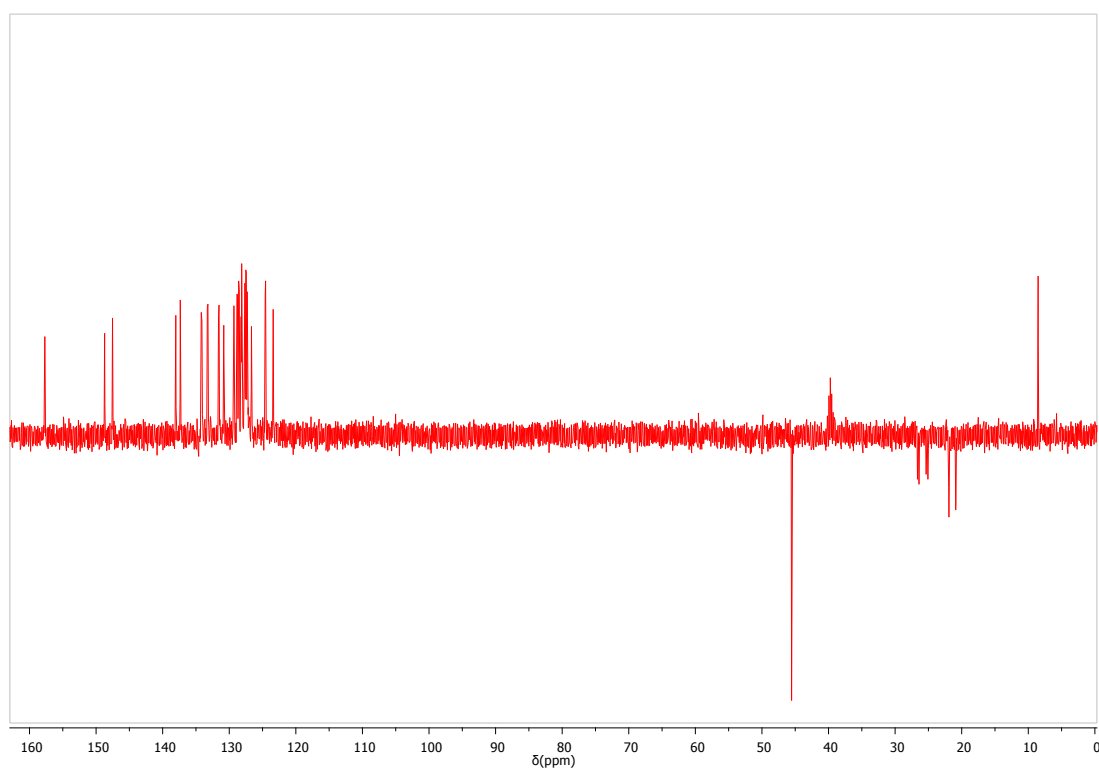


Figure S9 ^{13}C DEPT NMR spectra of the complex 1 (DMSO-d_6)

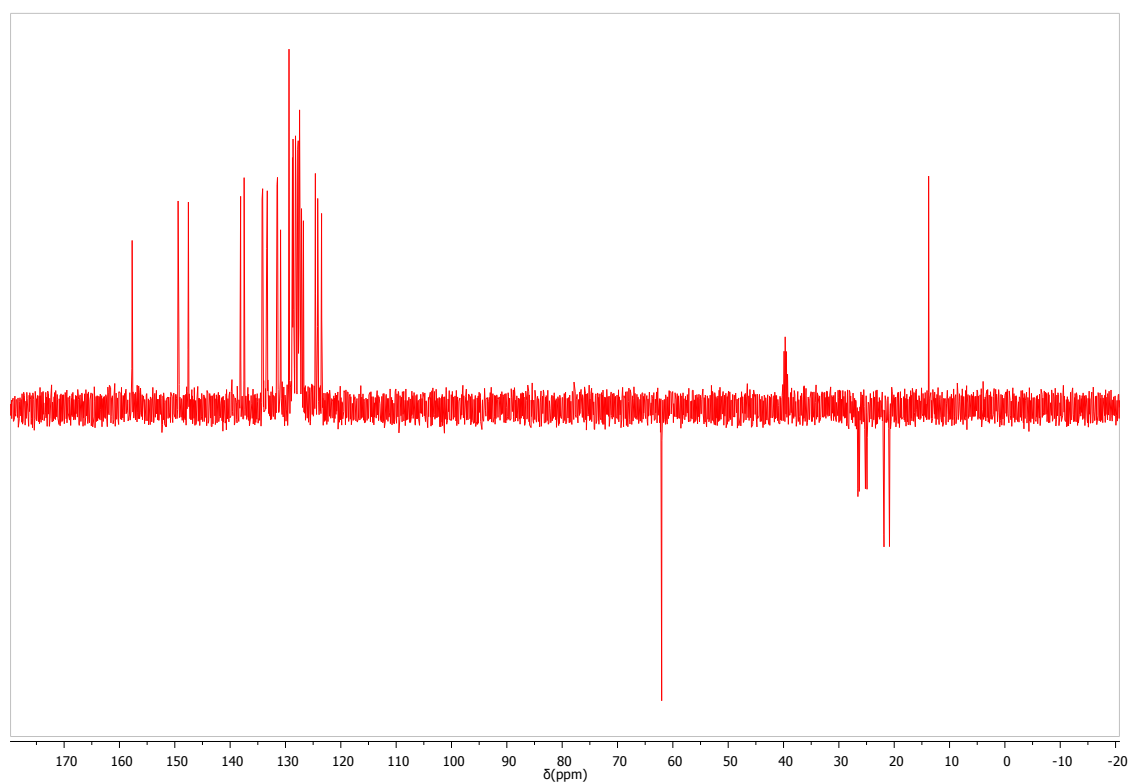


Figure S10 ^{13}C DEPT NMR spectra of the complex 2 (DMSO-d_6)

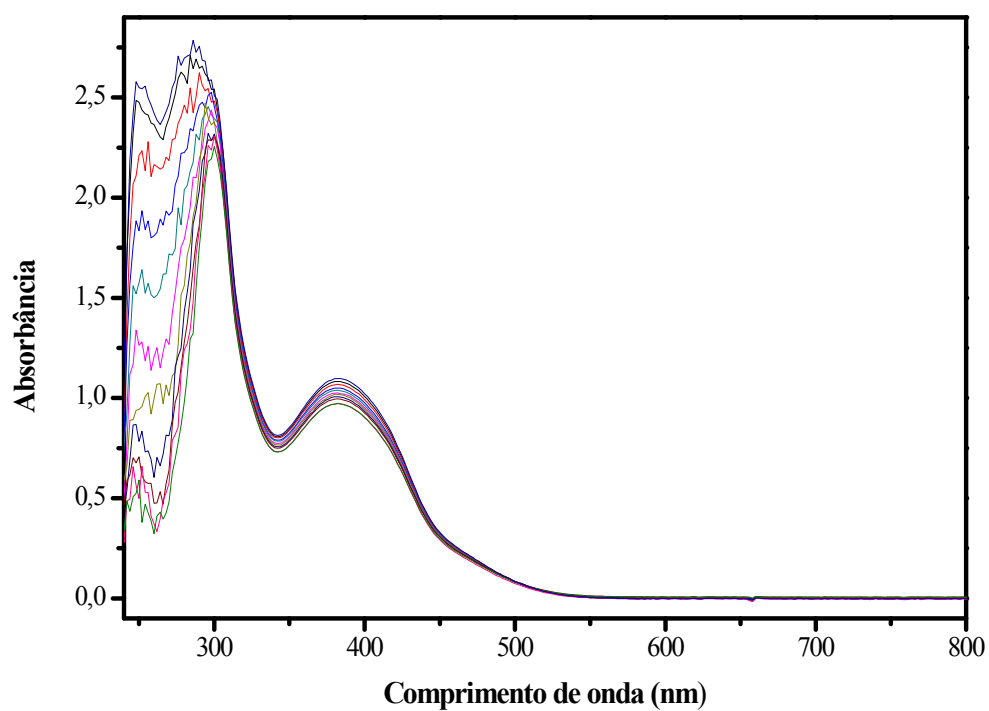


Figure S11 Spectrophotometric titration spectra of 1 with calf thymus (ct-DNA).

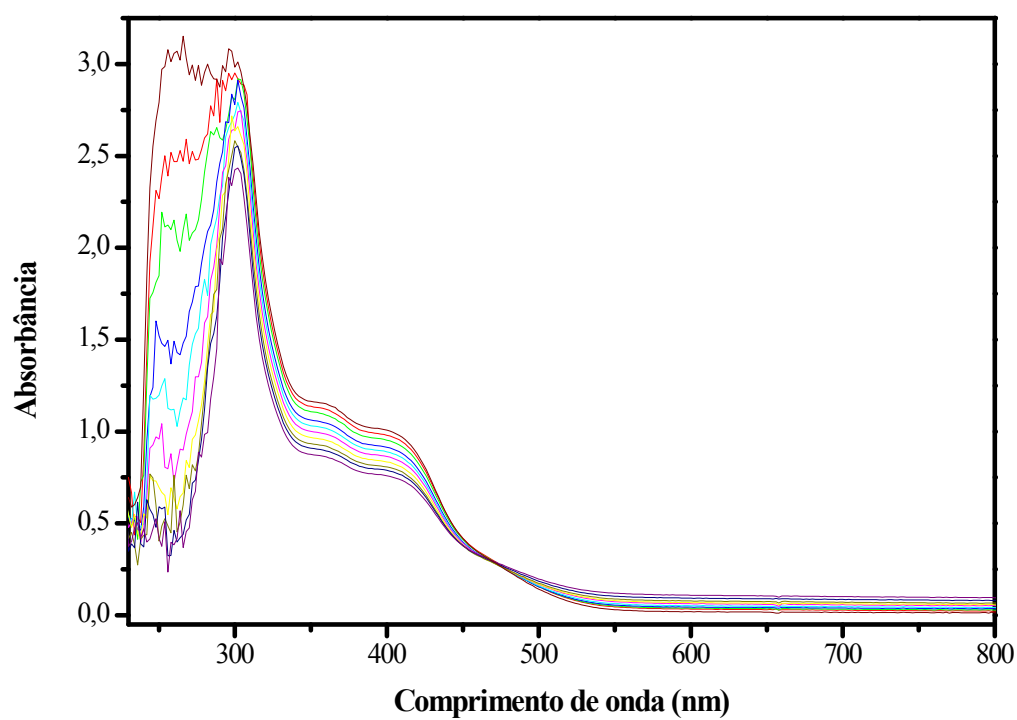


Figure S12 Spectrophotometric titration spectra of 3 with calf thymus (ct-DNA).

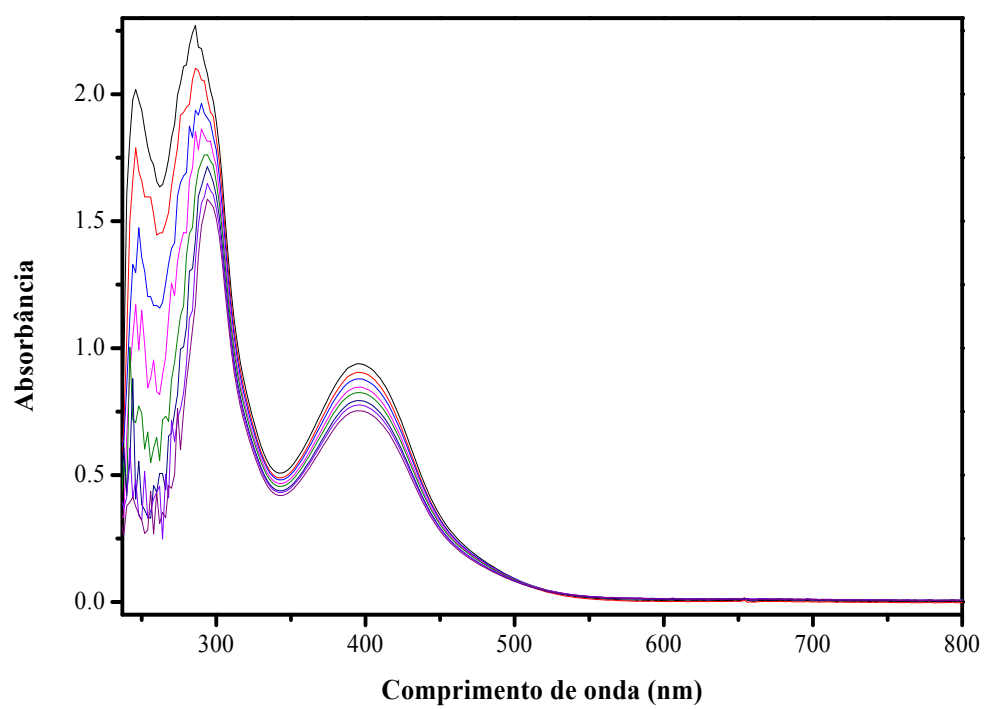


Figure S13 Spectrophotometric titration spectra of 2 with calf thymus (ct-DNA).

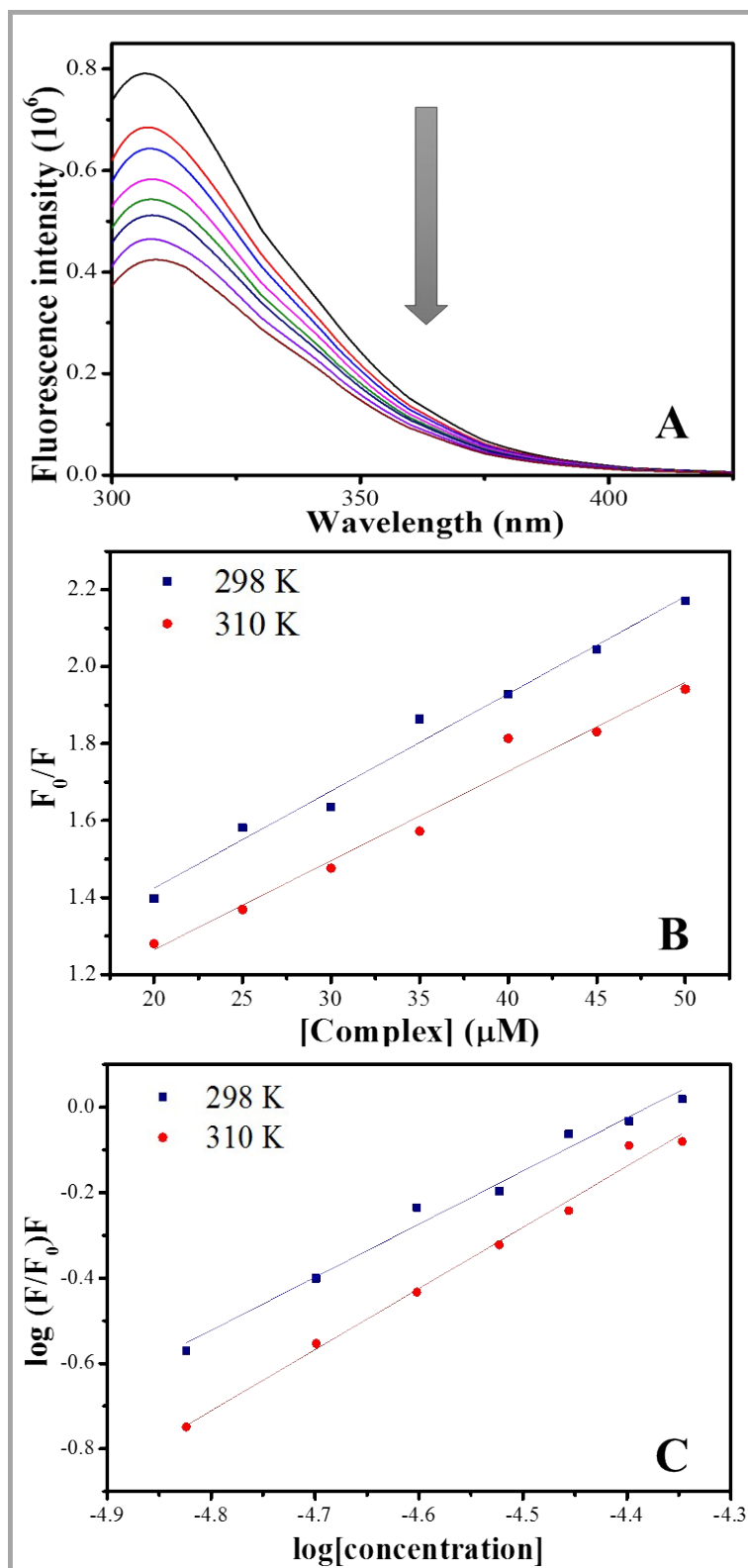


Figure S14 (A) Fluorescence quenching spectra of HSA with different concentrations of complex 2 in a Trizma-HCl buffer, pH 7.4; (B and C) Stern–Volmer plots for complex 2 showing tryptophan quenching at 310 and 289 K.

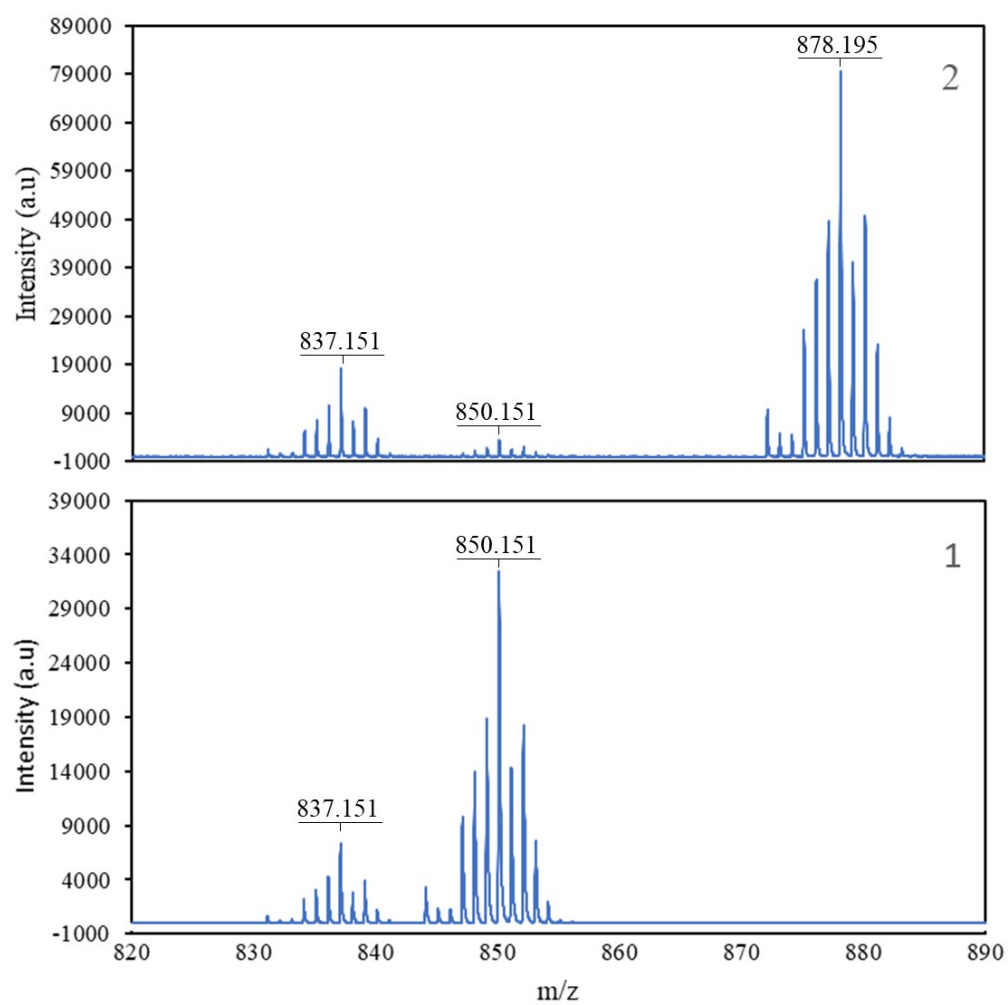


Figure S15 Maldi-TOF/MS spectra of complexes 1 and 2, using the dihydroxybenzoic acid as calibrant