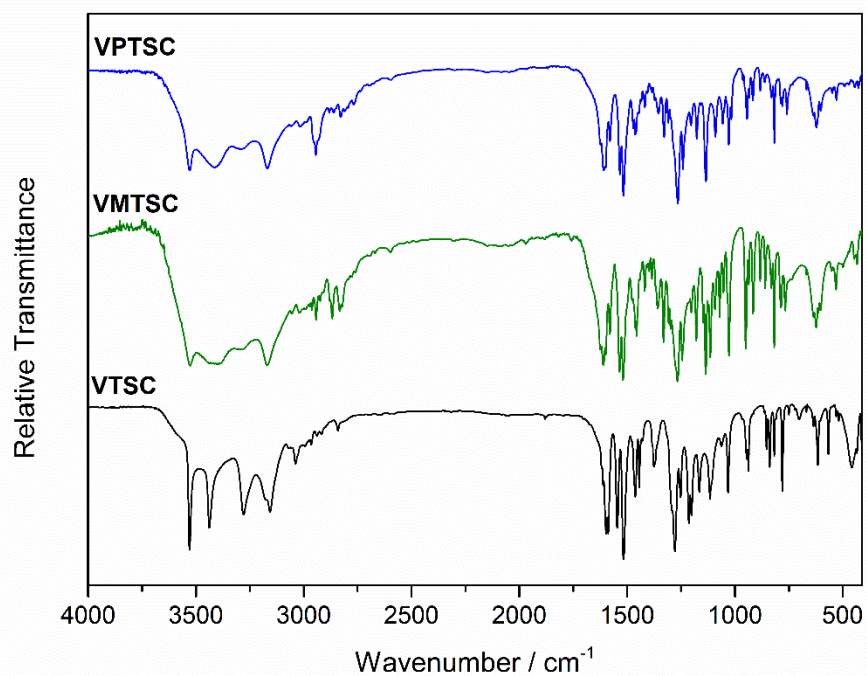


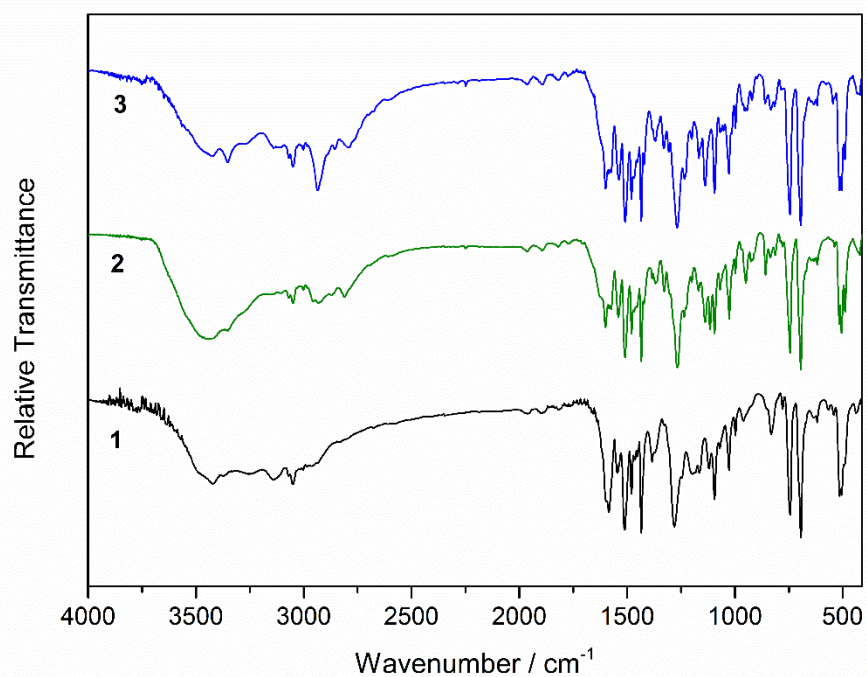
Electronic Supplementary Information (ESI)

Silver(I) complexes of 3-methoxy-4-hydroxybenzaldehyde thiosemicarbazones and triphenylphosphine: structural, cytotoxic and apoptotic studies

Débora E. S. Silva,^a  Amanda B. Becceneri,^b João V. B. Santiago,^a José A. Gomes Neto,^a Javier Ellena,^c Márcia R. Cominetti,^b José C. M. Pereira,^a Michael J. Hannon^d and Adelino V. G. Netto^a 



(A)

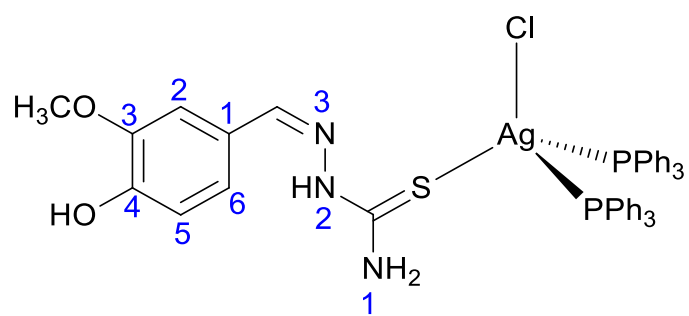


(B)

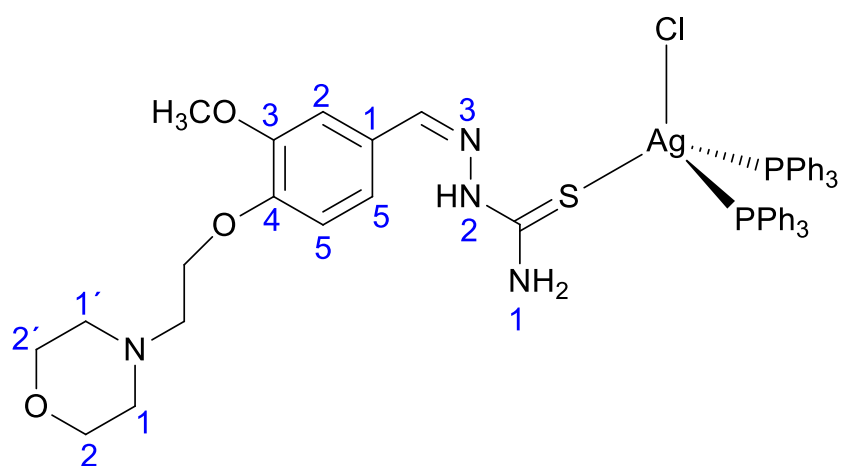
Fig. S1. IR spectra of (A) thiosemicarbazones and (B) complexes **1-3**.

Table S1 Selected bond lengths (Å) and bond angles (°) of complexes **2** and **3**.

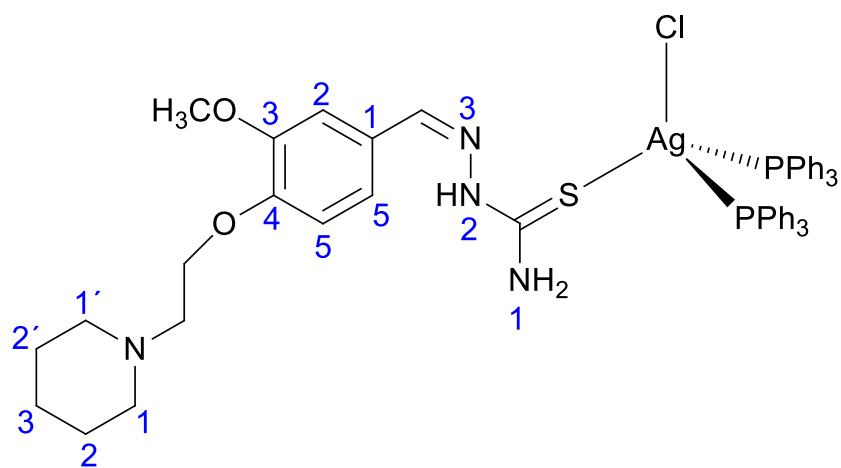
2	Length (Å)		Angle (°)
Ag(1) – S(21)	2.6013(7)	P(31) – Ag(1) – S(21)	107.06(2)
Ag(1) – P(31)	2.4917(7)	P(31) – Ag(1) – Cl(44)	103.89(3)
Ag(1) – Cl(44)	2.5892(8)	Cl(44) – Ag(1) – S(21)	107.86(2)
Ag(1) – P(51)	2.4759(7)	P(51) – Ag(1) – S(21)	109.04(2)
		P(51) – Ag(1) – P(31)	115.35(3)
		P(51) – Ag(1) – Cl(44)	113.21(3)
3			
Ag(1) – S(1)	2.6325(8)	P(8) – Ag(1) – Cl(40)	100.72(2)
Ag(1) – P(8)	2.4794(7)	P8 – Ag(1) – S(1)	114.40(3)
Ag(1) – Cl(40)	2.5723(7)	P(27) – Ag(1) – P(8)	124.21(2)
Ag(1) – P(27)	2.4661(6)	P(27) – Ag(1) – Cl(40)	113.22(2)
		P(27) – Ag(1) – S(1)	99.80(2)
		Cl(40) – Ag(1) – S(1)	103.09(2)



(1)

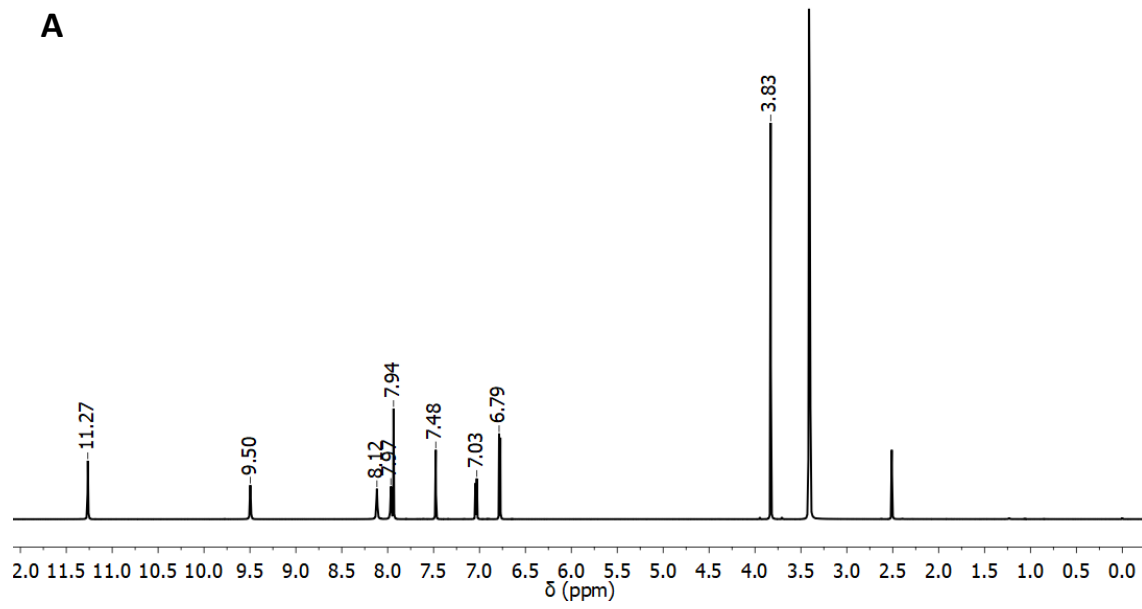
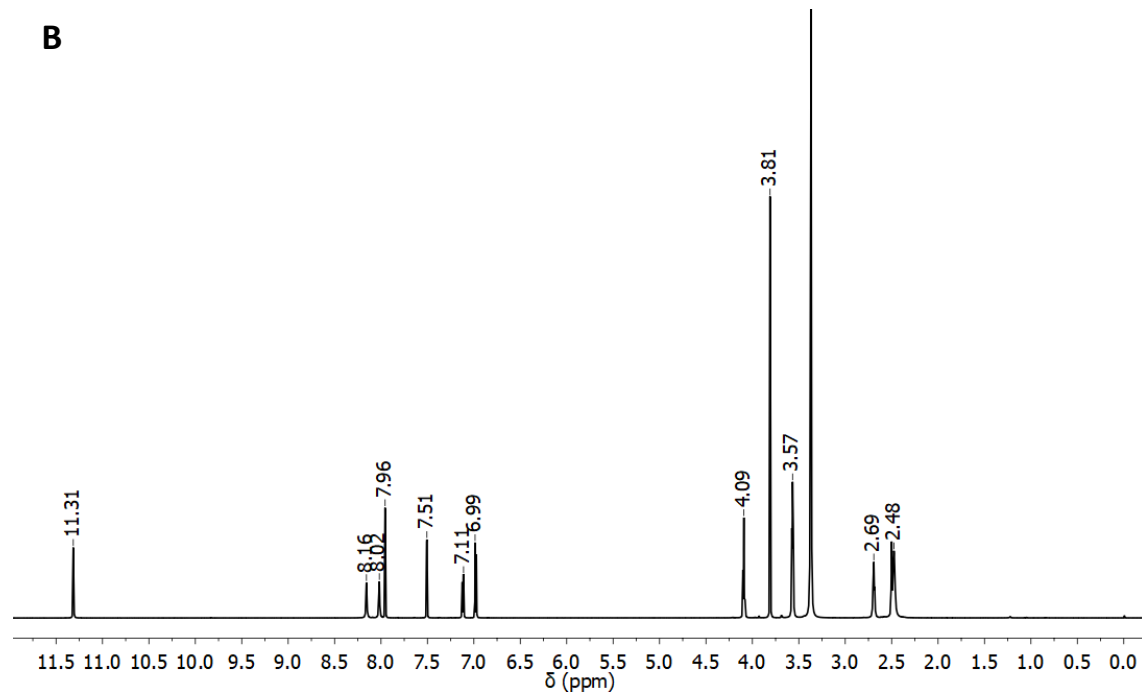


(2)



(3)

Fig. S2. Numbering scheme of complexes **1-3** for NMR data.

A**B**

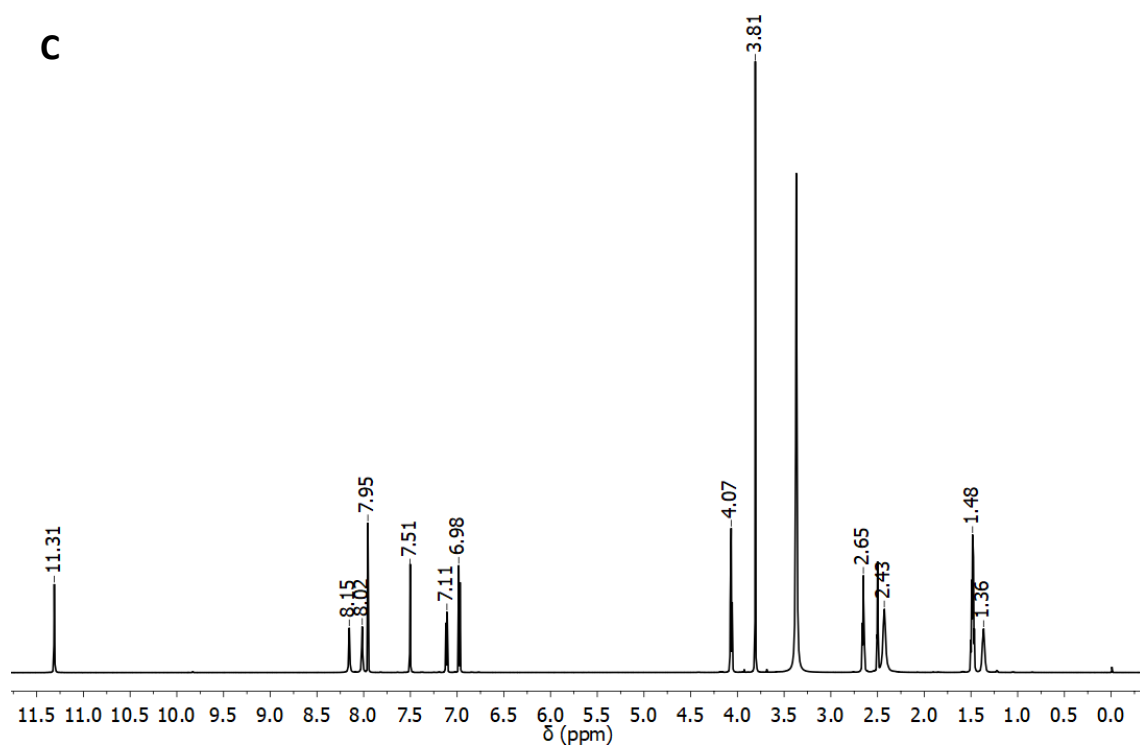
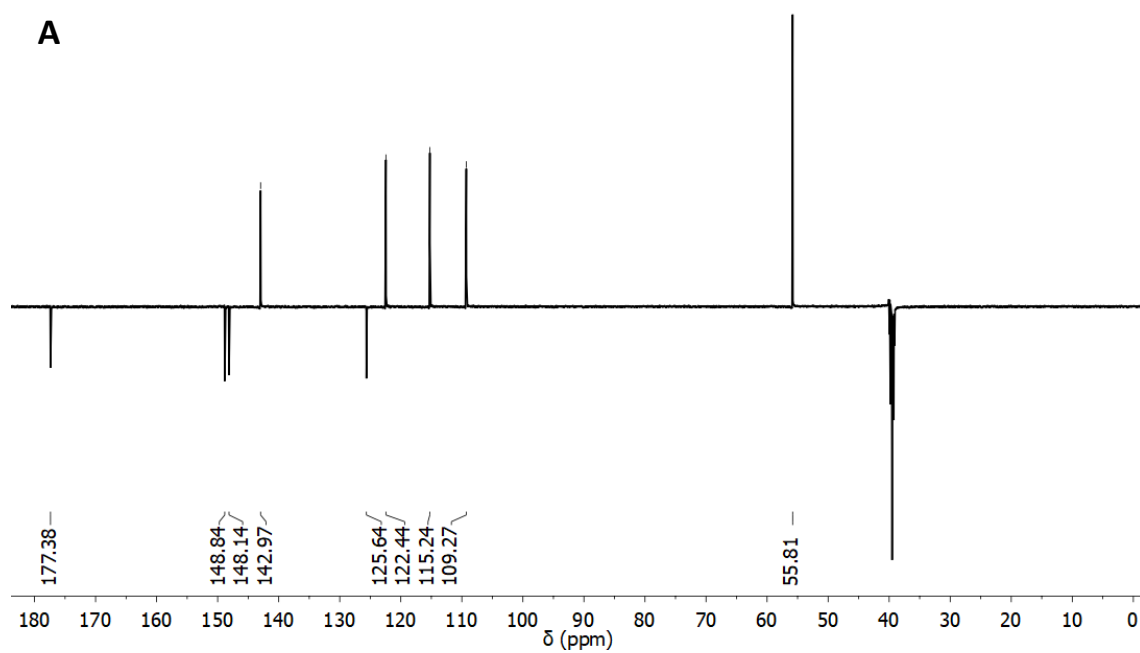


Fig. S3. ^1H NMR spectra of thiosemicabazones (A) VTSC; (B) VMTSC and (C) VPTSC in DMSO-d_6 .



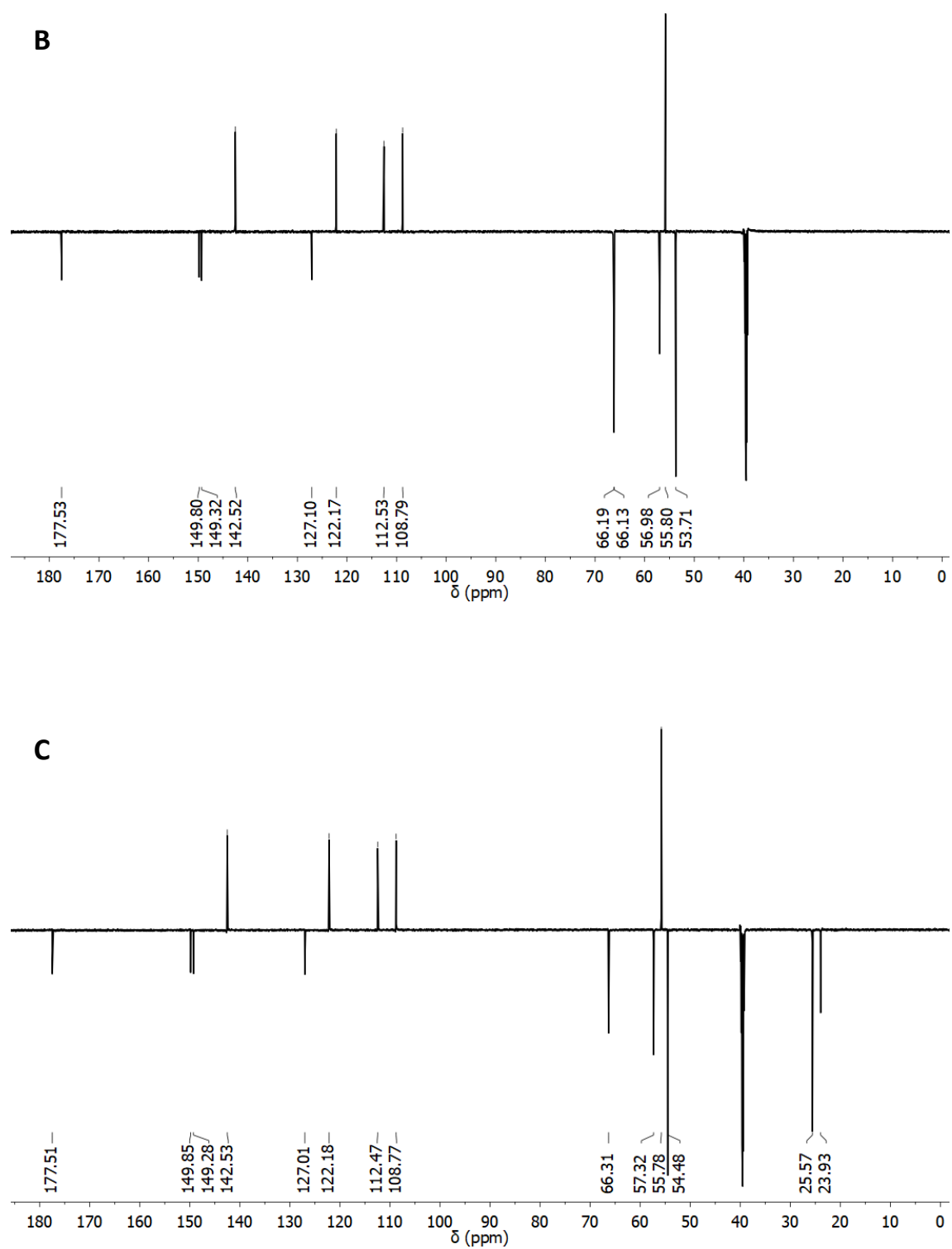


Fig. S4. DEPTQ NMR spectra of thiosemicabazones (A) VTSC; (B) VMTSC and (C) VPTSC in DMSO- d_6 .

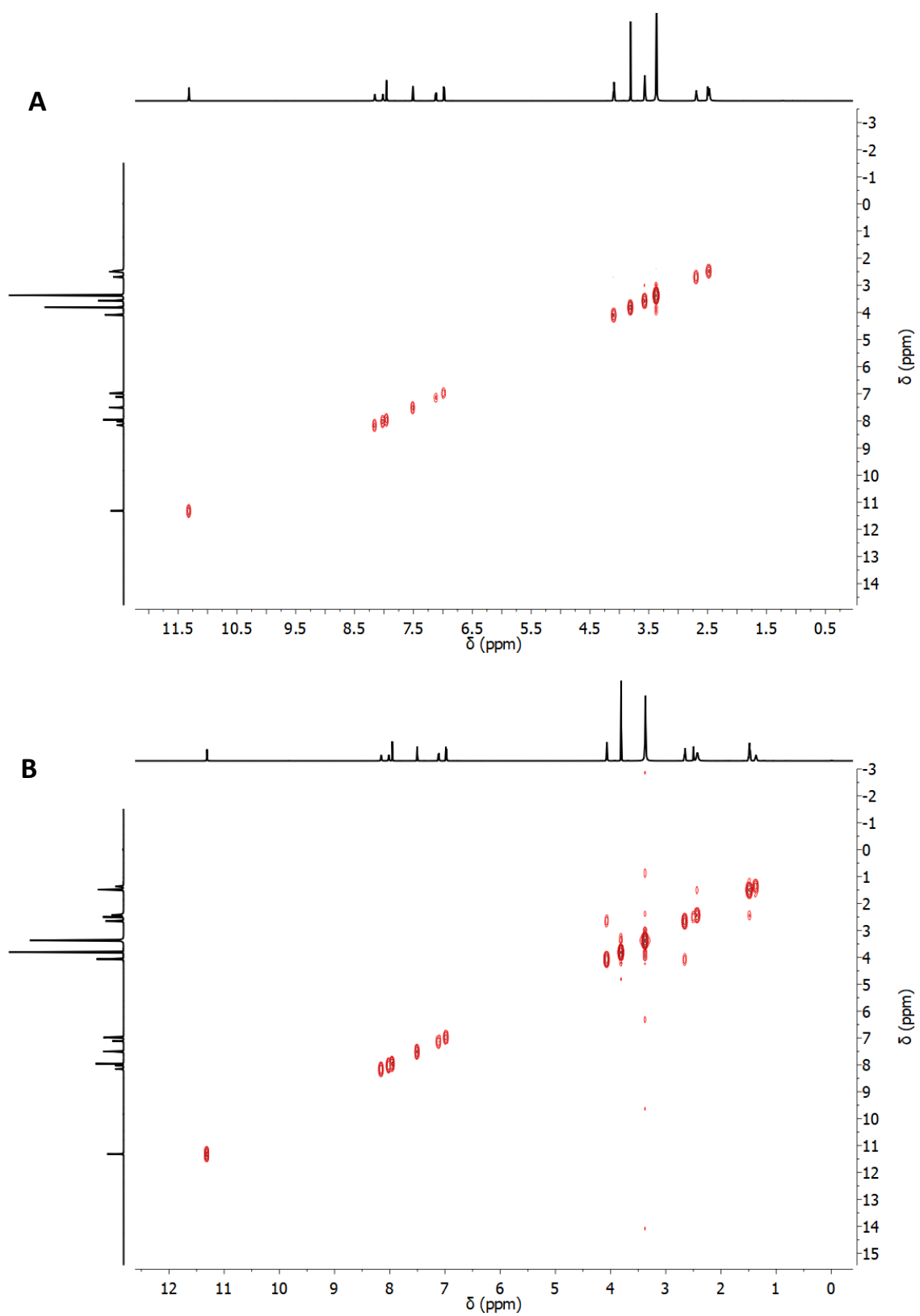
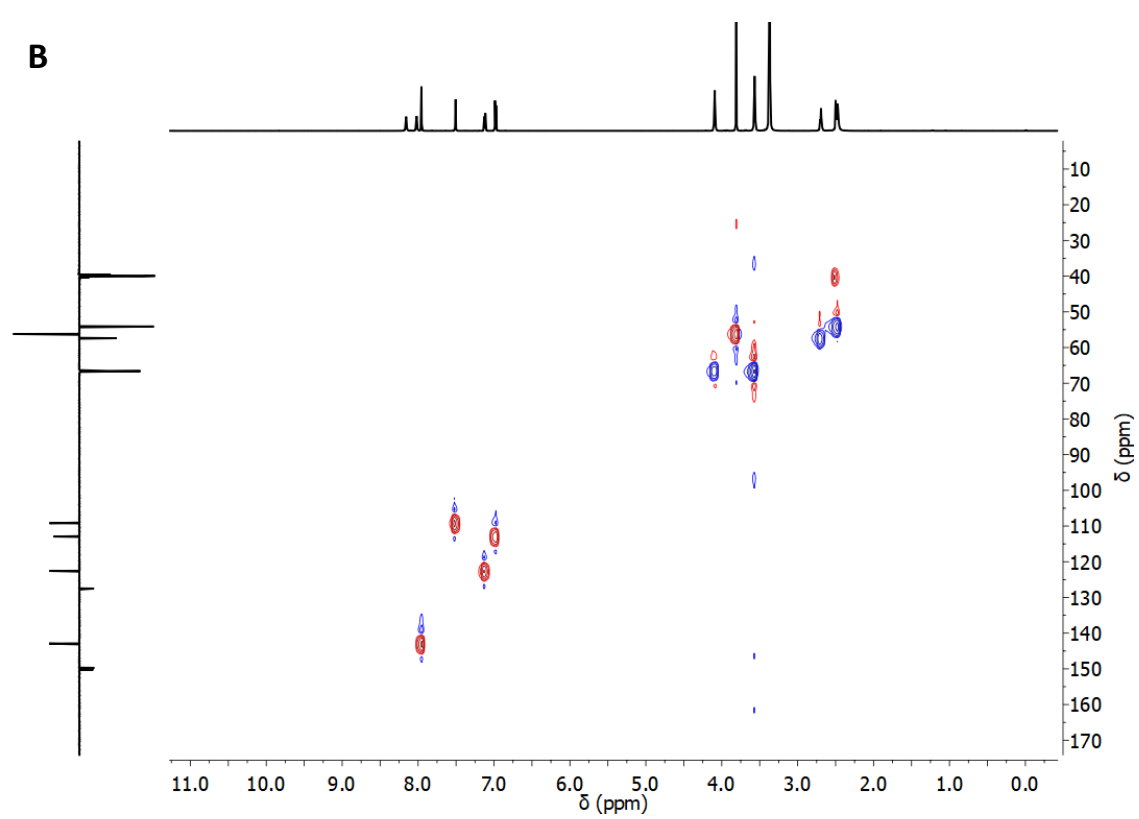
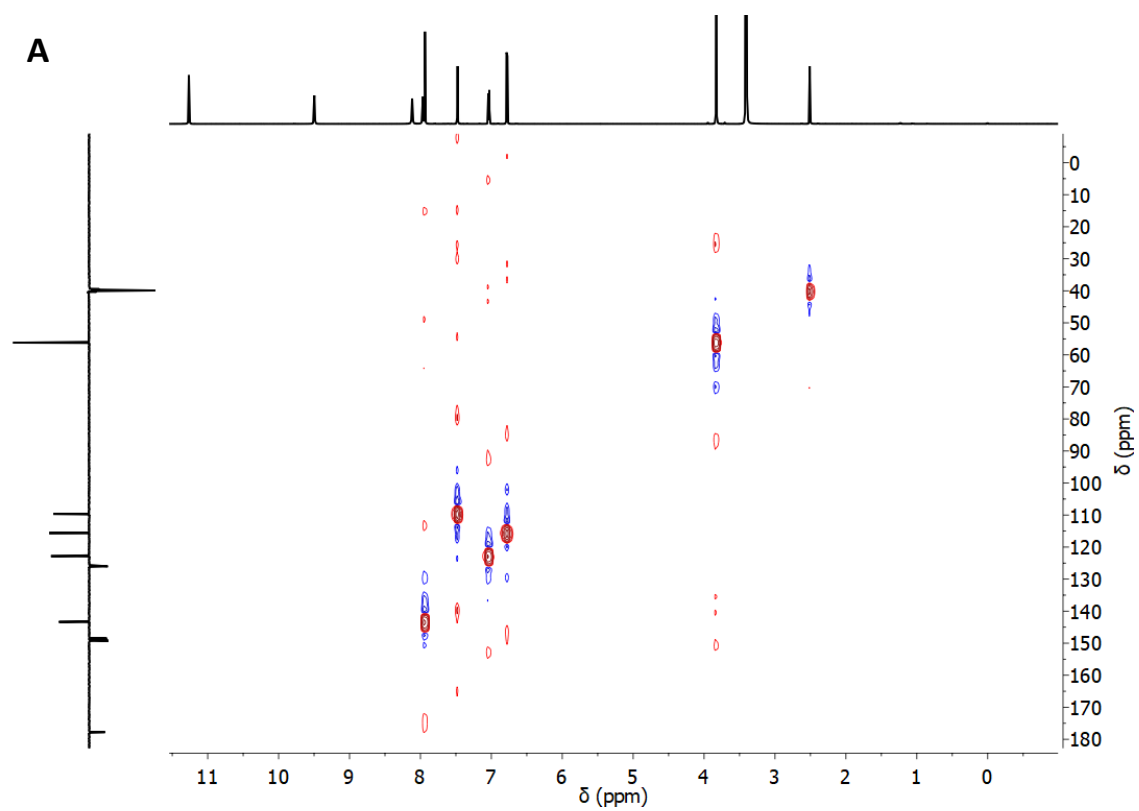


Fig. S5. COSY ^1H - ^1H NMR spectra of thiosemicabazones (A) VMTSC and (B) VPTSC in DMSO-d_6 .



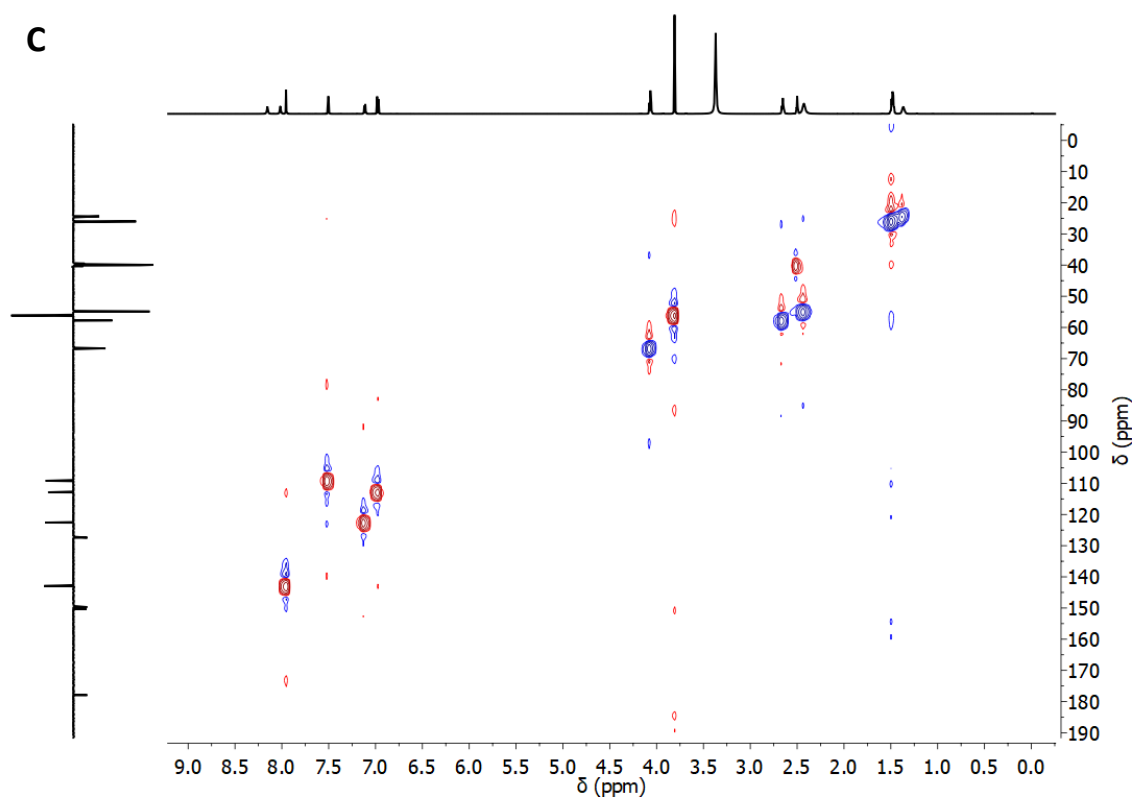
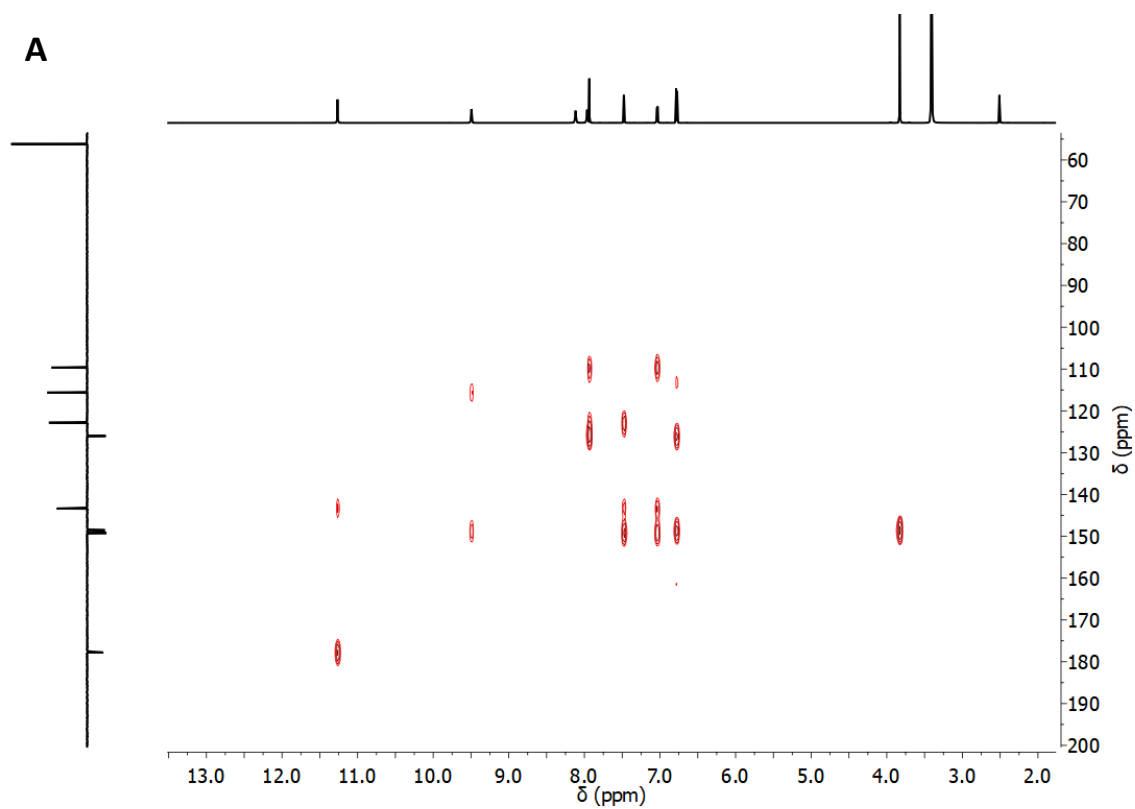


Fig. S6. ^1H - ^{13}C HSQC spectra of thiosemicabazones (A) VTSC; (B) VMTSC and (C) VPTSC in DMSO-d_6 .



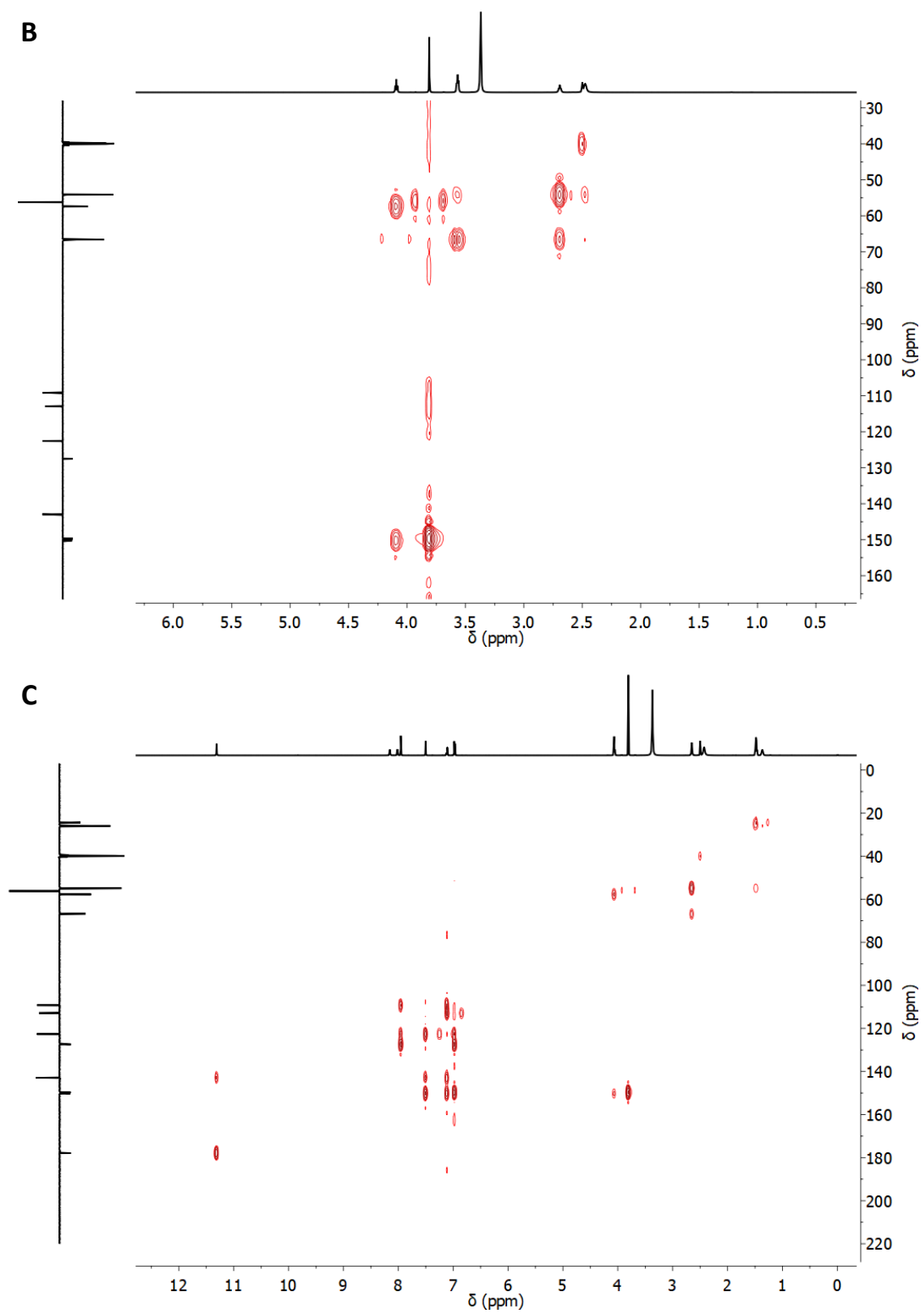
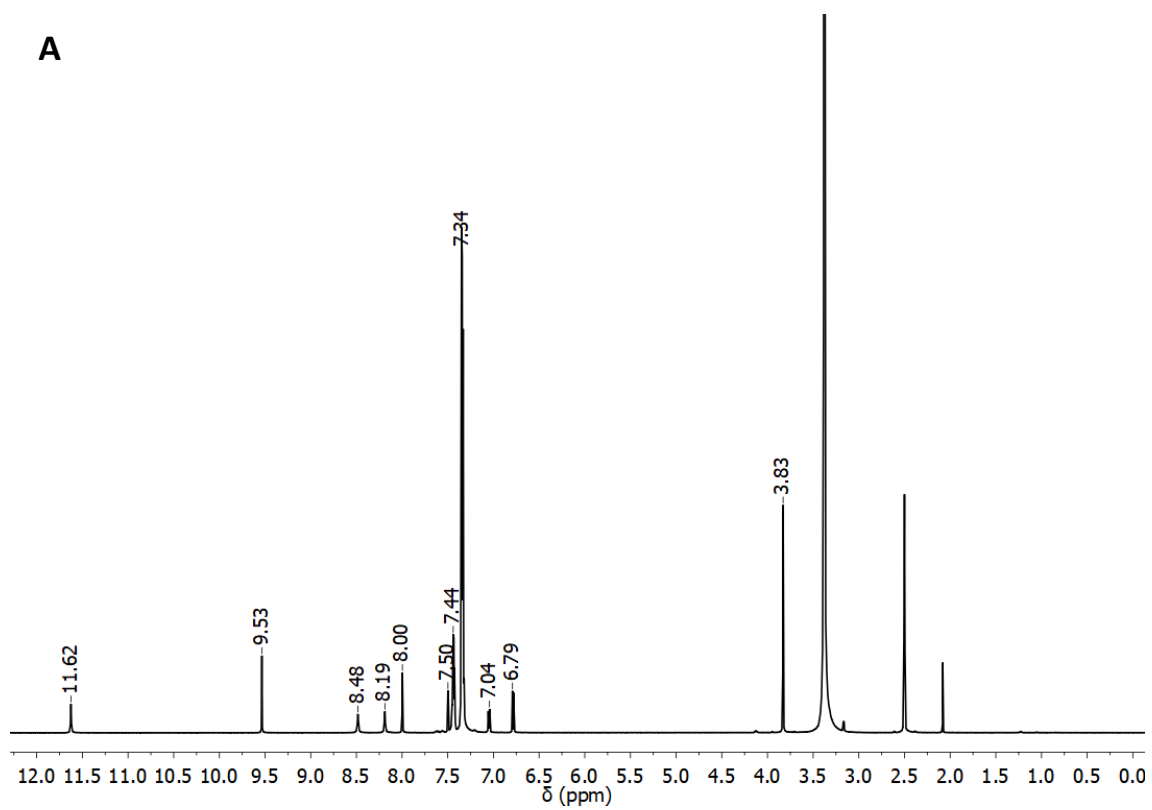
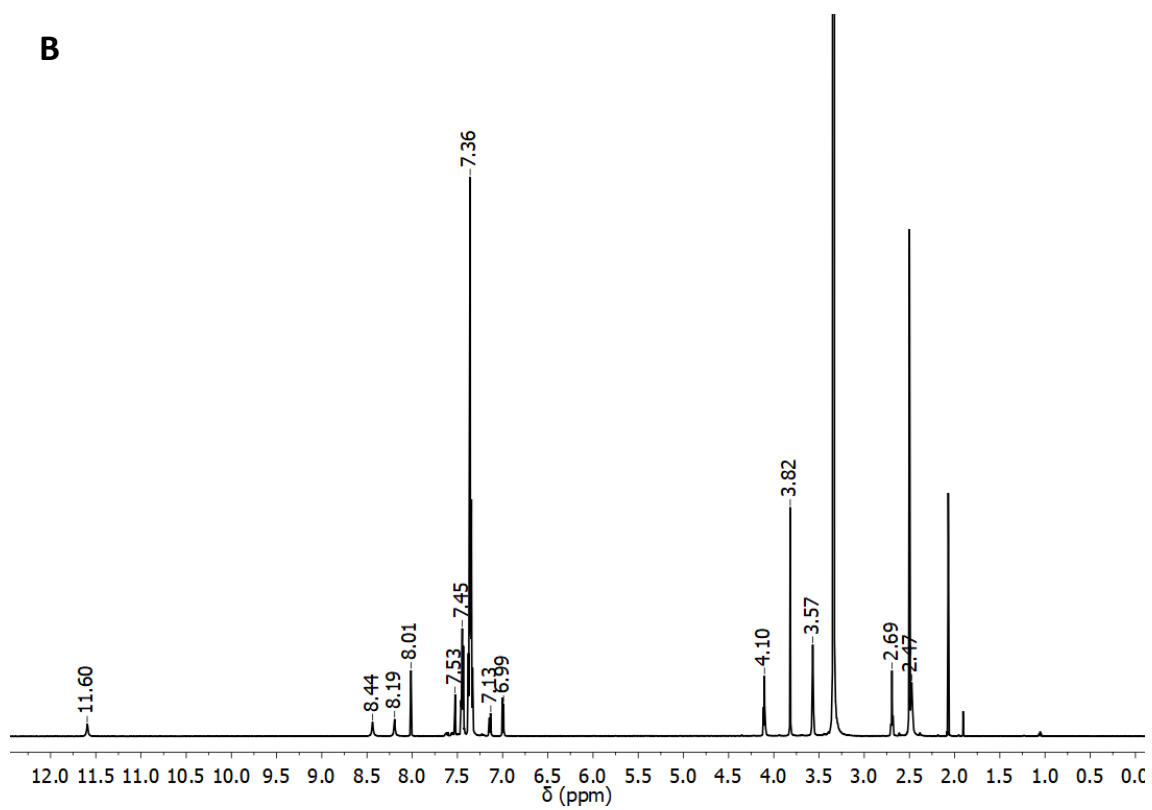


Fig. S7. ^1H - ^{13}C HMBC spectra of thiosemicabazones (A) VTSC; (B) VMTSC and (C) VPTSC in DMSO-d_6 .

A**B**

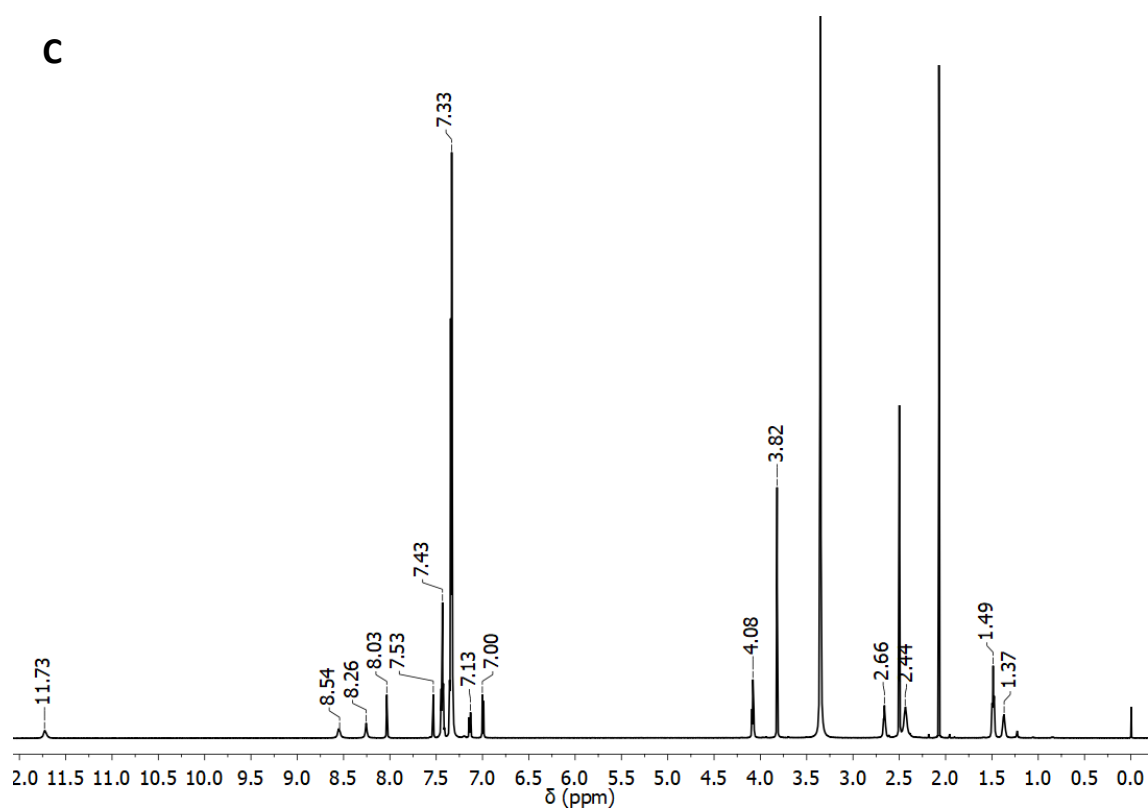
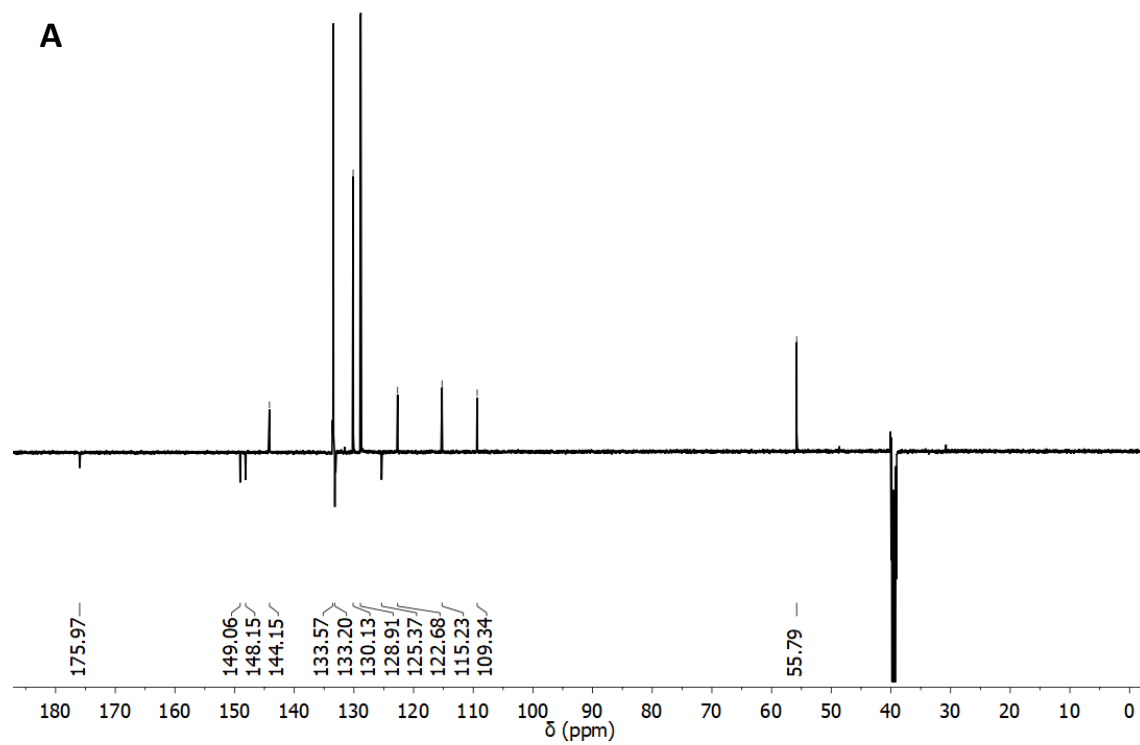


Fig. S8. ^1H NMR spectra of complexes (A) **1**; (B) **2** and (C) **3**.



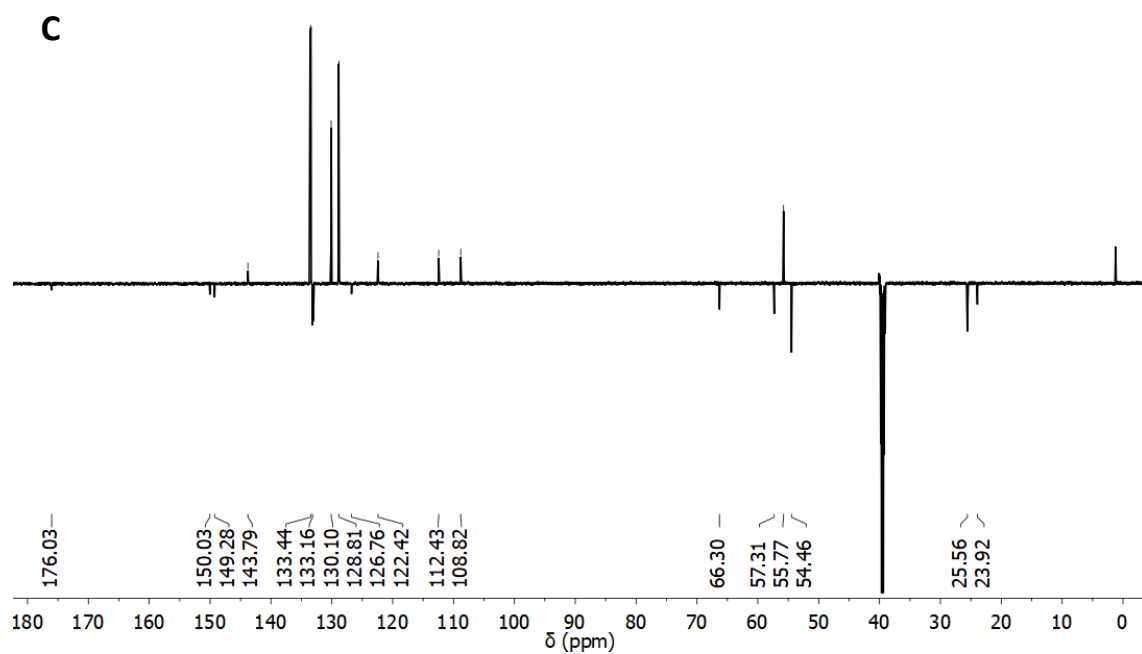
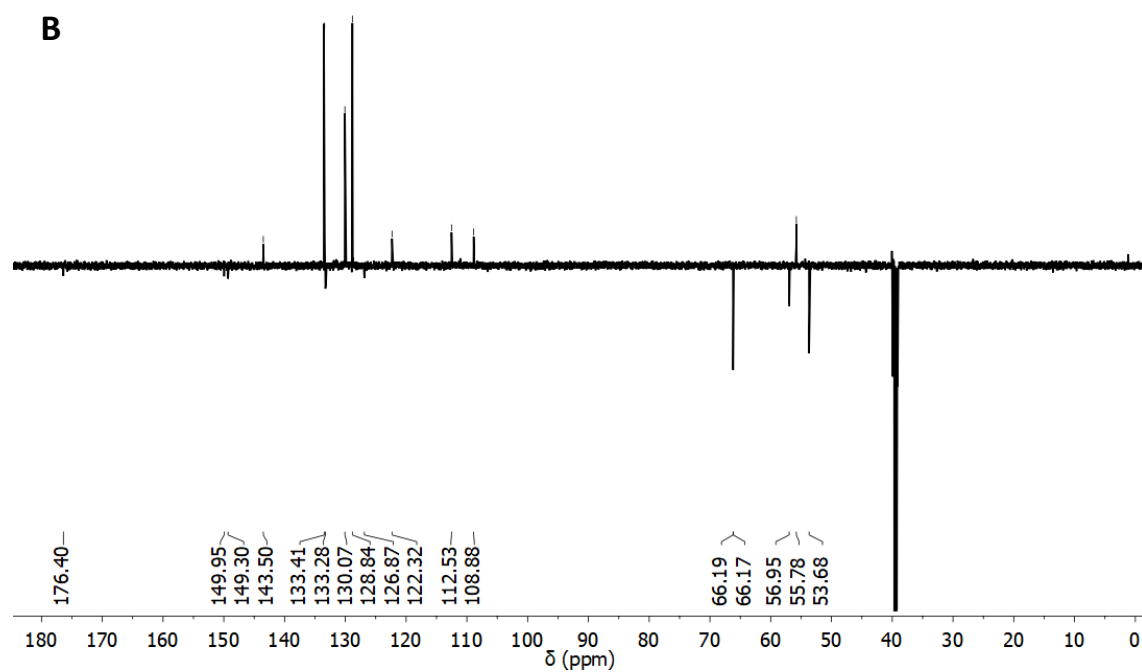
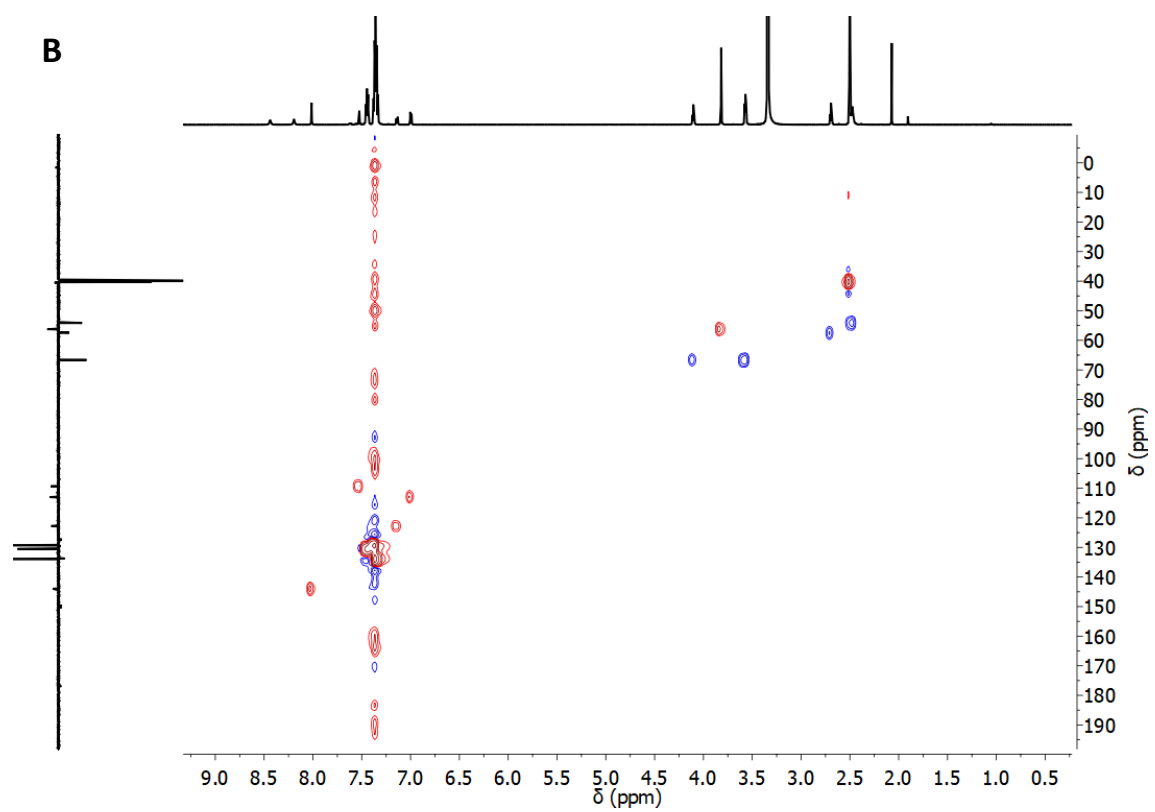
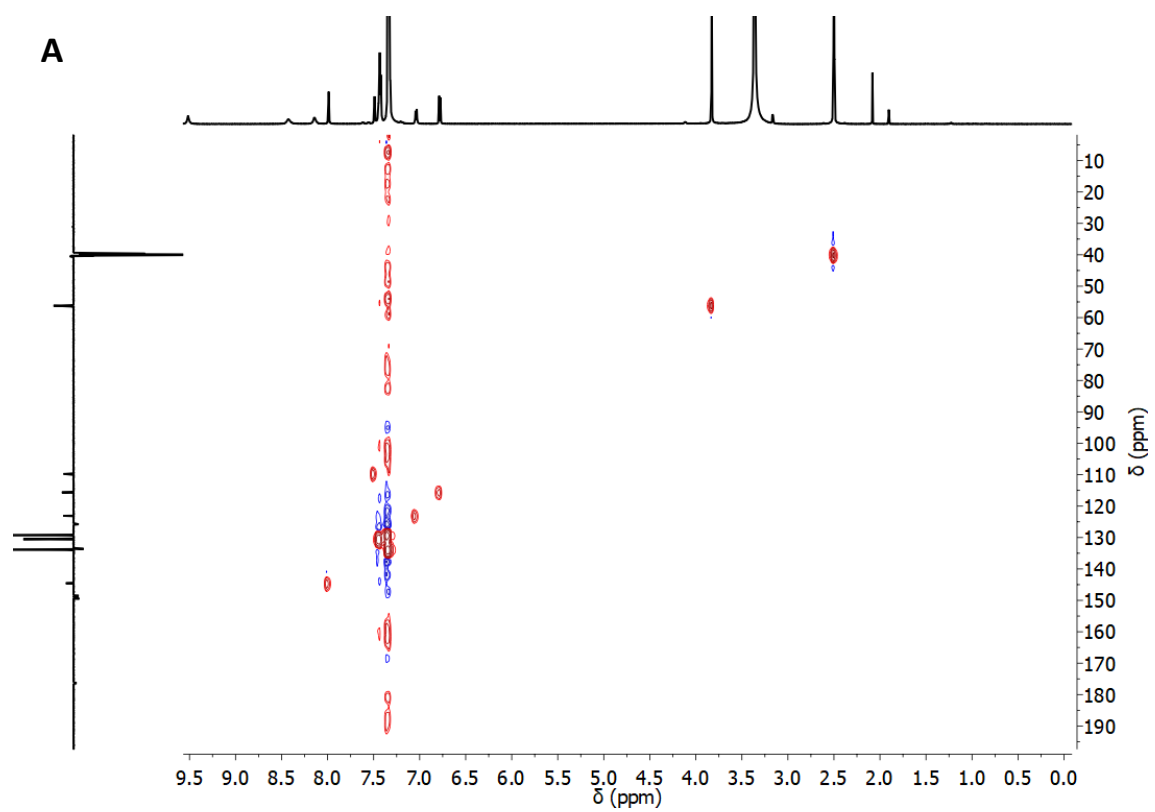


Fig. S9. DEPTQ NMR spectra of complexes (A) **1**; (B) **2** and (C) **3** in DMSO- d_6 .



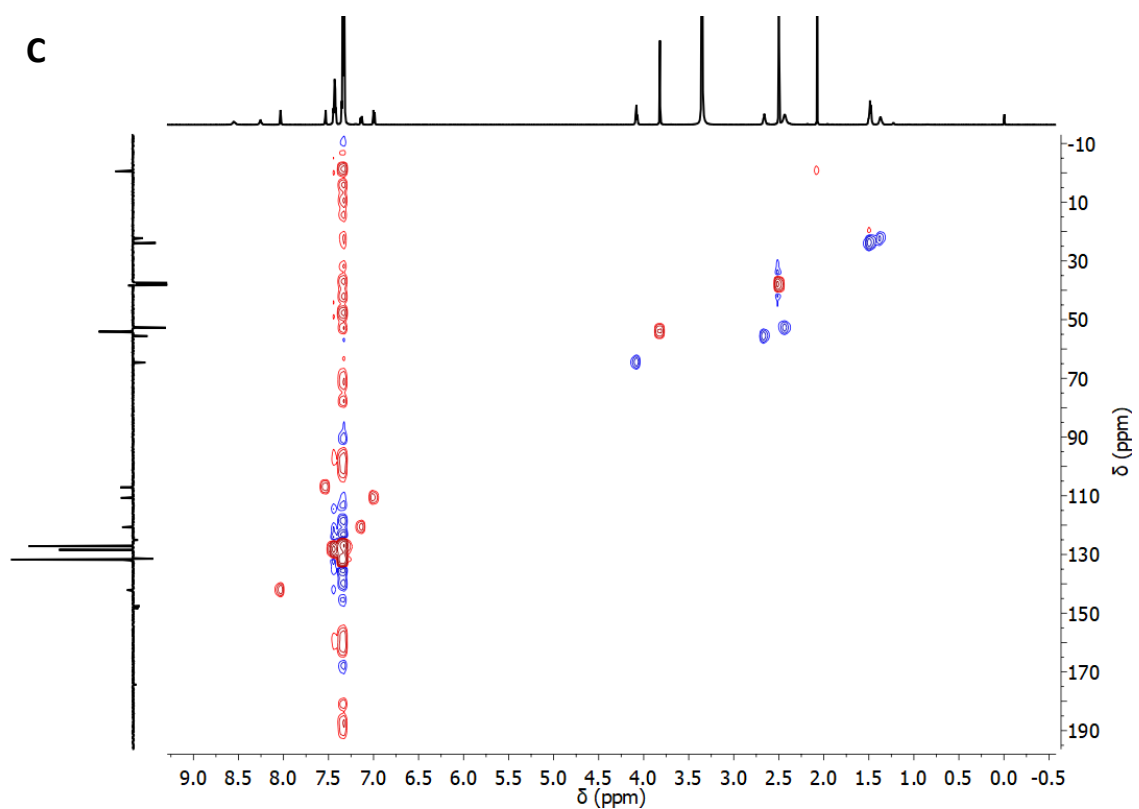
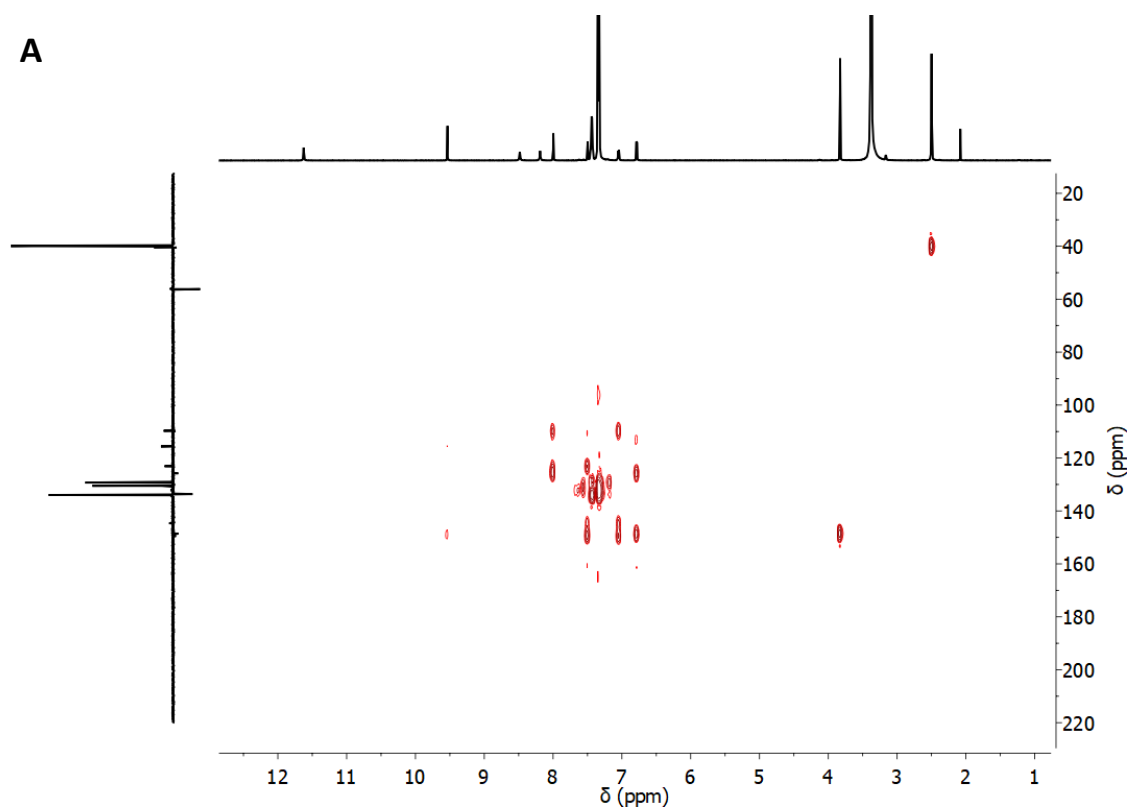


Fig. S10. ^1H - ^{13}C HSQC spectra of complexes (A) **1**; (B) **2** and (C) **3** in DMSO-d_6 .



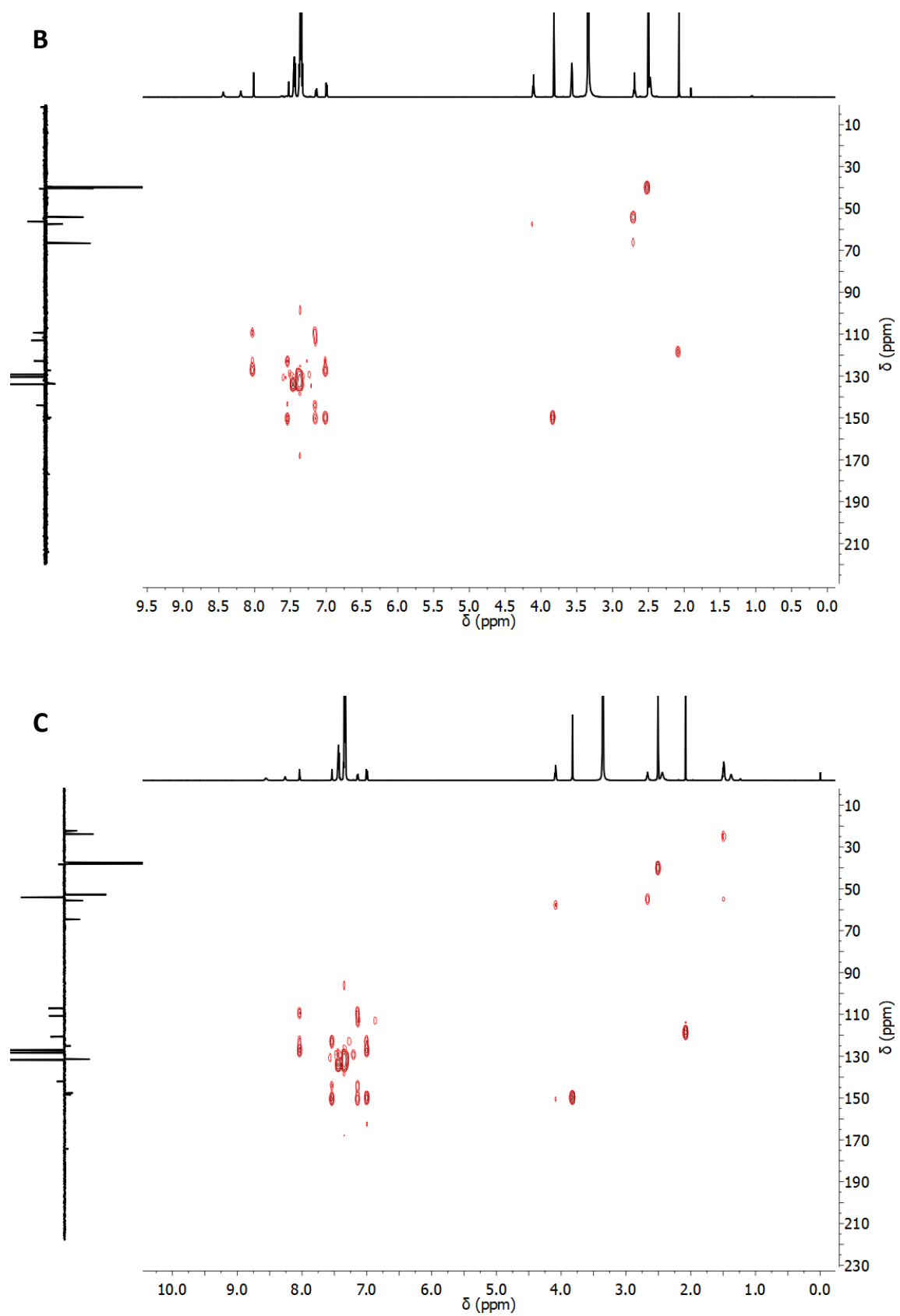


Fig. S11. ^1H - ^{13}C HMBC spectra of complexes (A) **1**; (B) **2** and (C) **3** in DMSO-d_6 .

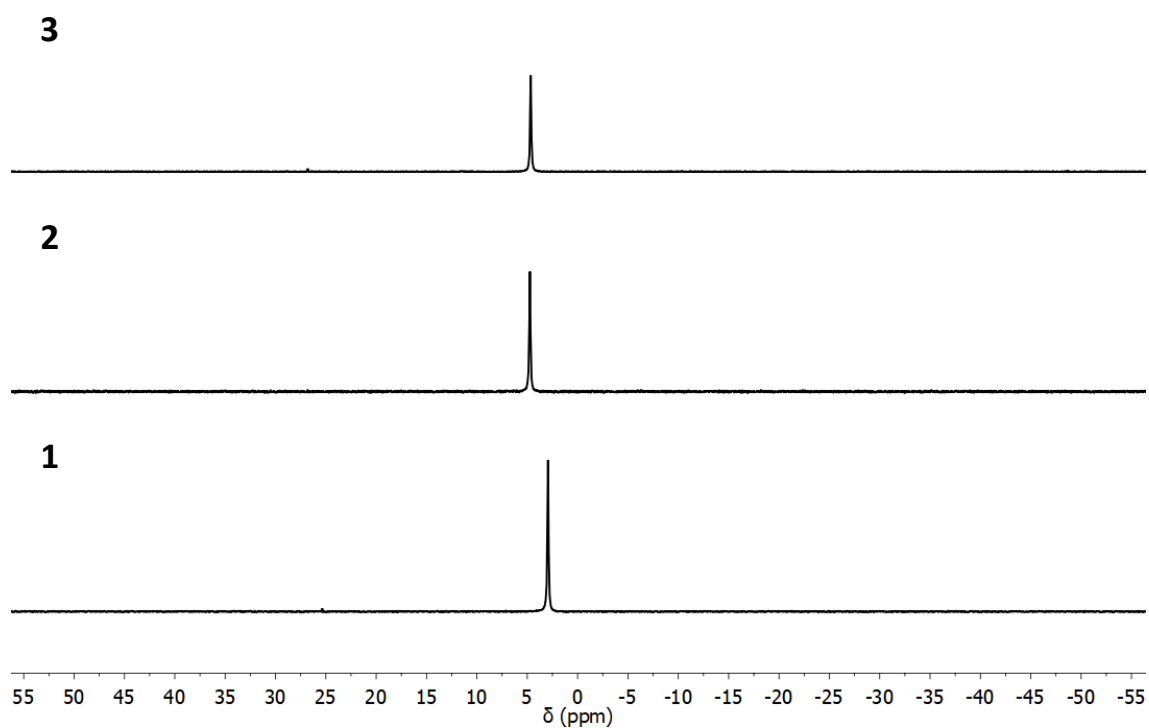
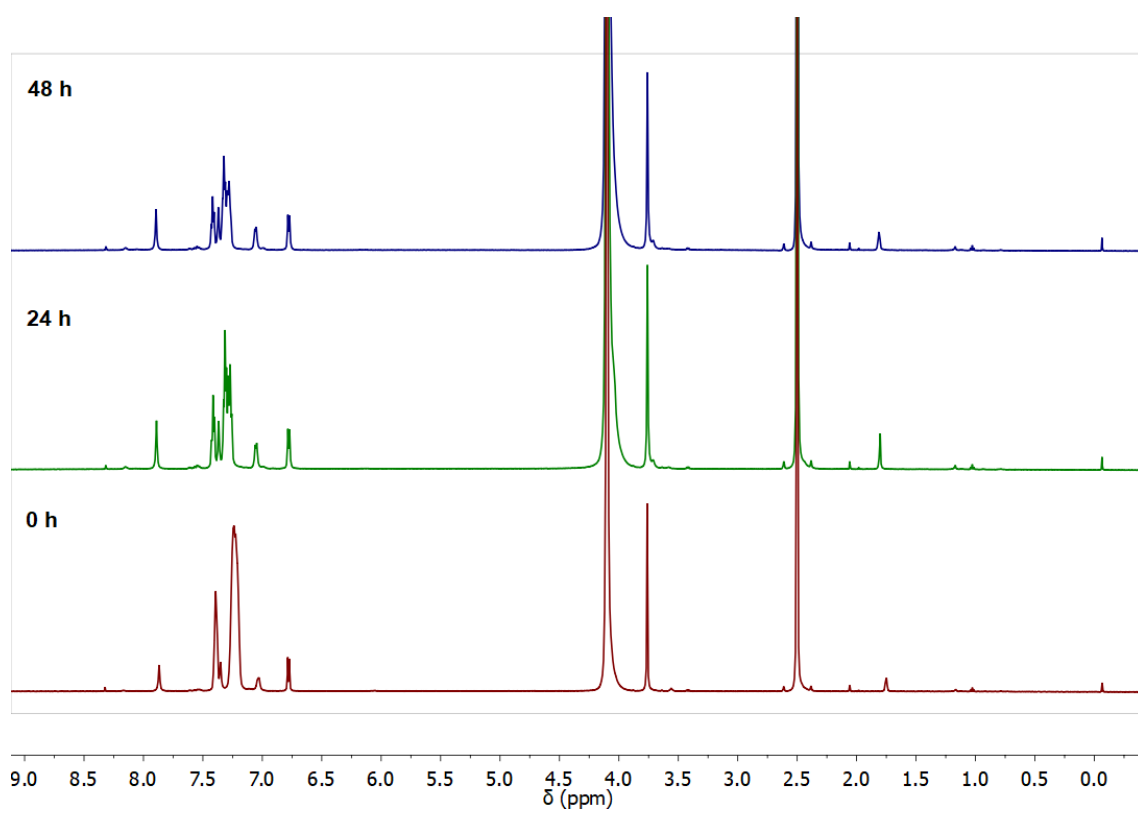
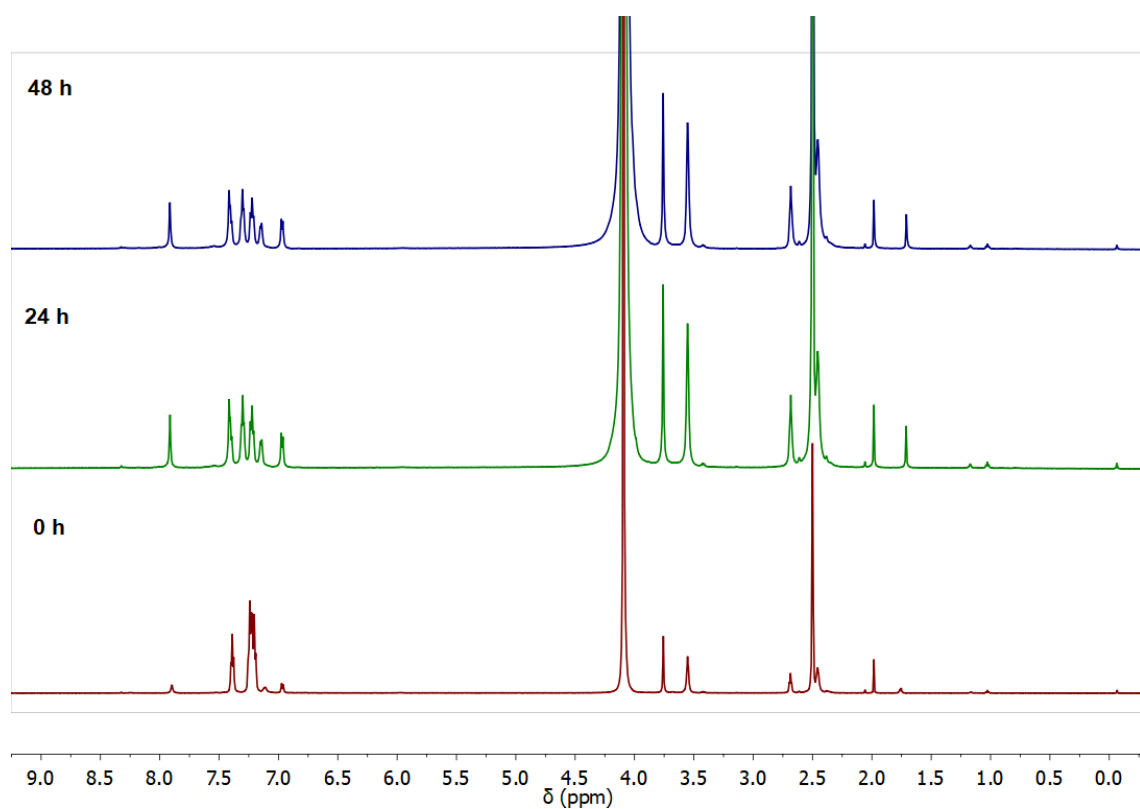


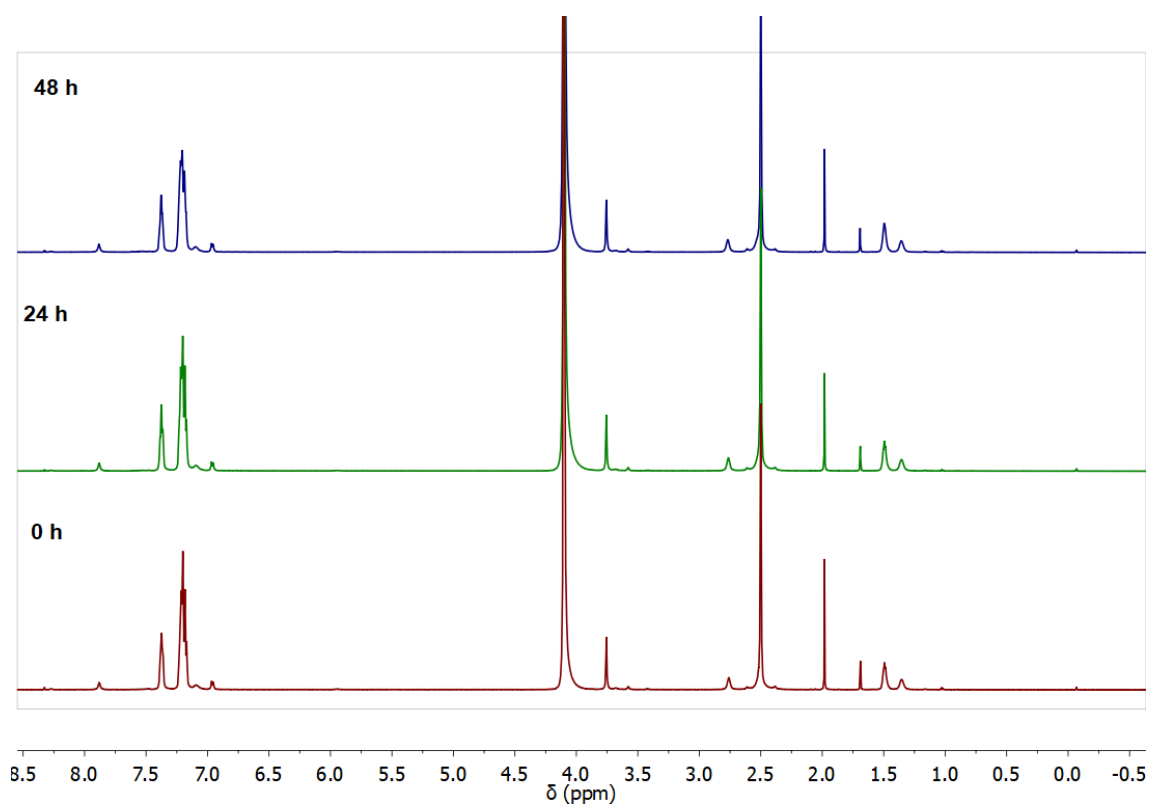
Fig. S12. ^{31}P NMR spectra of complexes **1-3** in DMSO-d_6 .



(A)



(B)



(C)

Fig. S13. ^1H NMR spectra of complexes (A) **1**, (B) **2** and (C) **3** recorded in DMSO- d_6 /D $_2$ O (80; 20% v/v) over a period of 48 hours.

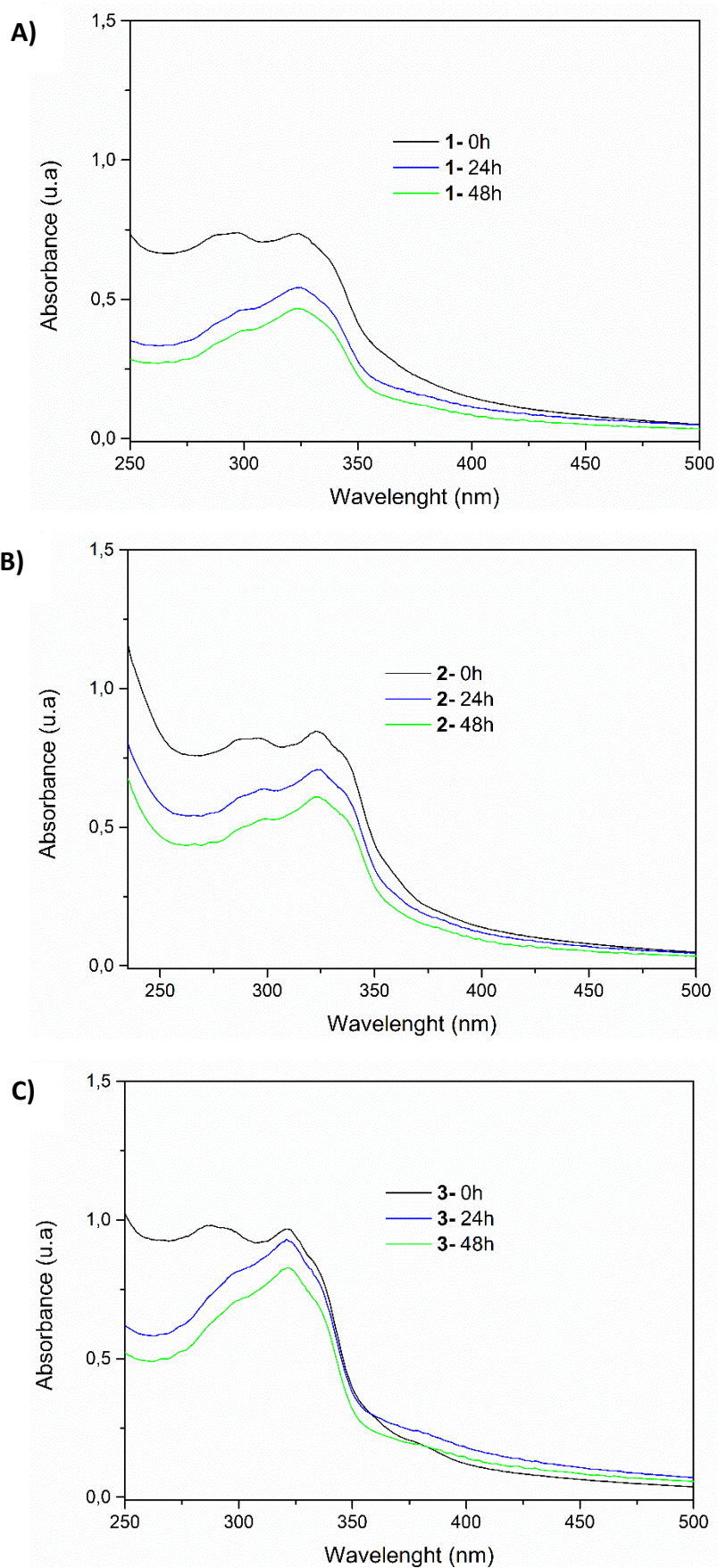


Fig. S14. UV-Vis spectra of complexes (A) **1**, (B) **2** and (C) **3** in Tris-HCl buffer (2% DMSO) recorded at 0, 24h and 48h after preparation of solutions.

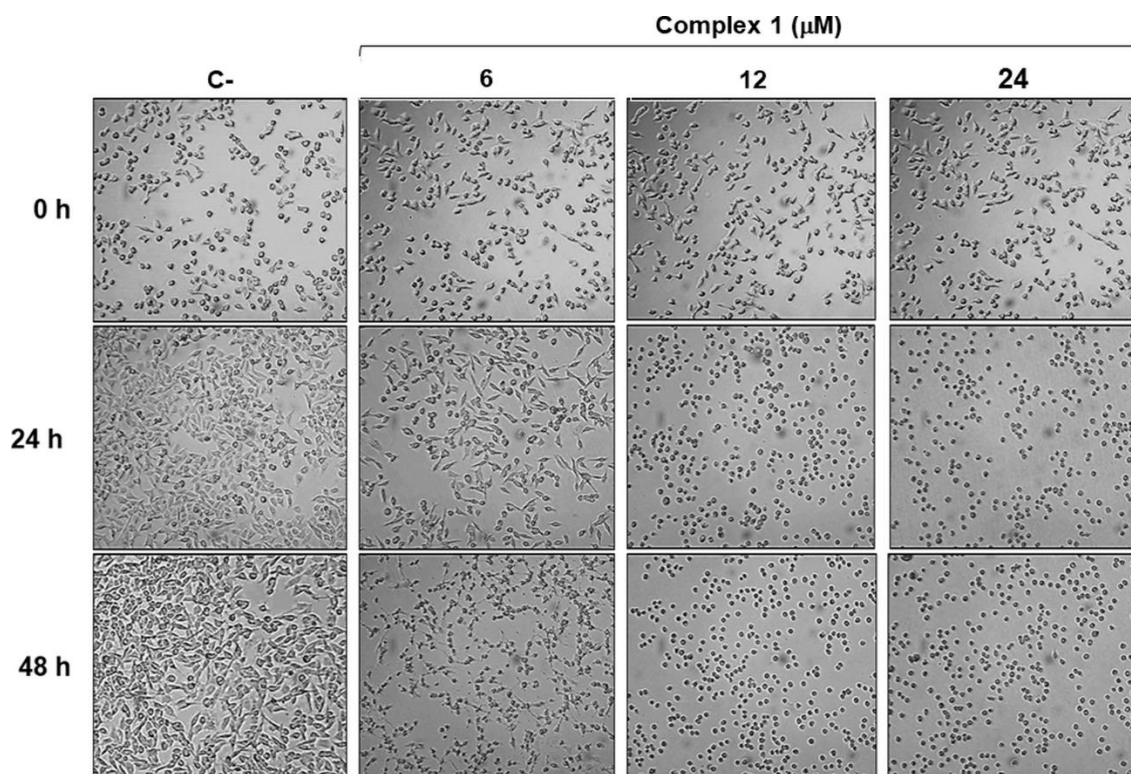


Fig. S15. Morphological changes in MDA-MB-231 cells incubated with complex **1** at a period of 48 hours. Amplification 10 x.

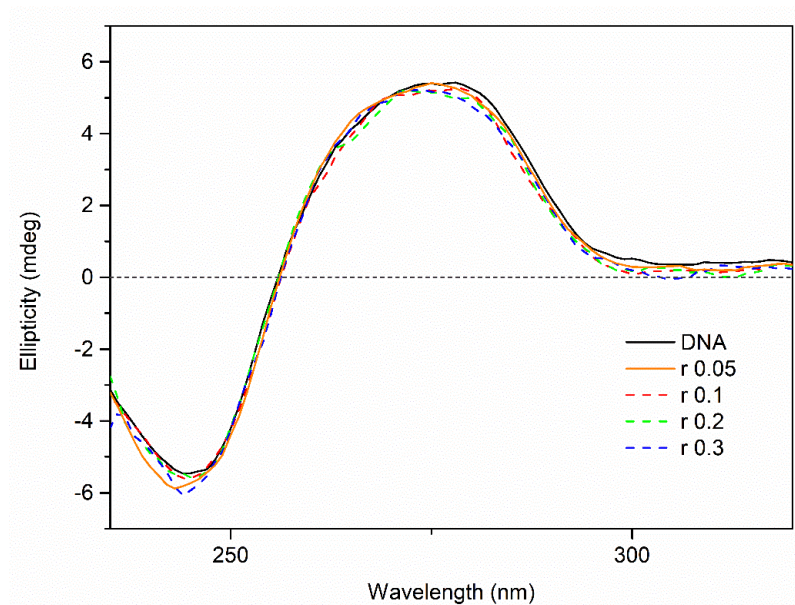


Fig. S16. CD spectrum of ct-DNA in the presence of **1**. $r = [\text{complex } 1] / [\text{DNA}]$.

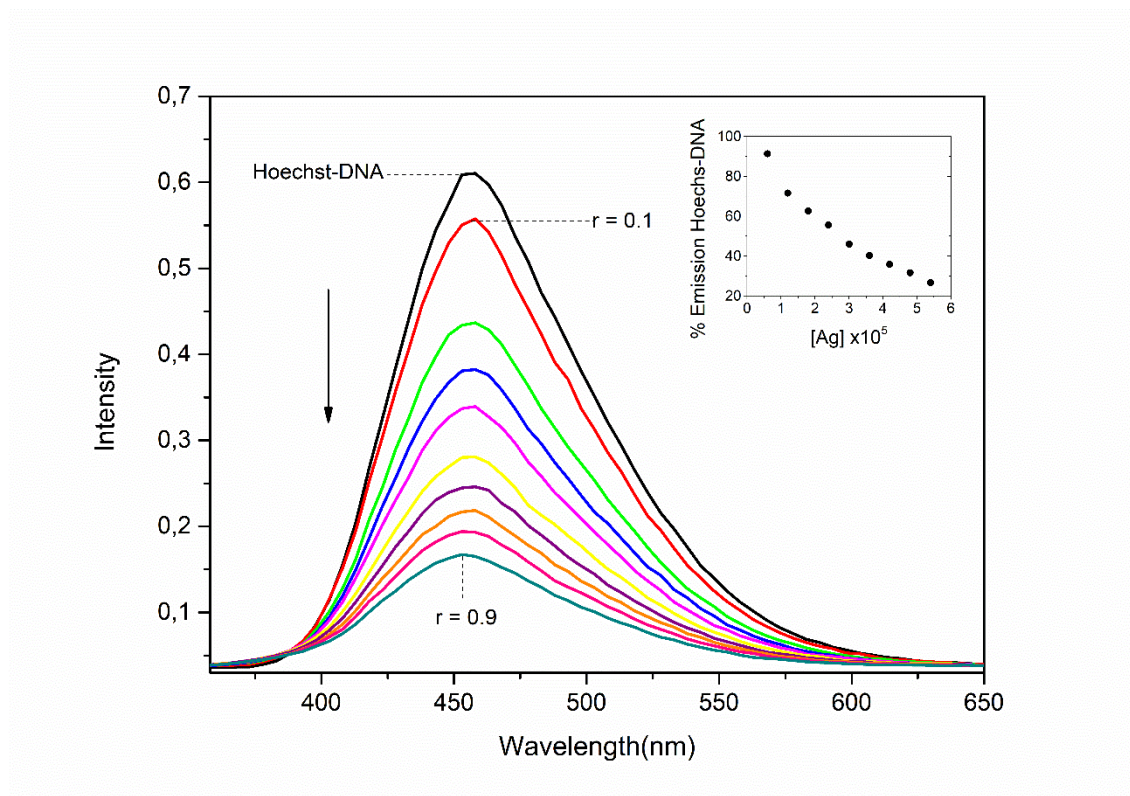


Fig S17. Emission spectra of Hoechst-bound DNA ($\lambda_{\text{ex}} = 350 \text{ nm}$) in the presence of increasing concentrations of **1** (6 - 60 μM) in Tris-HCl buffer. [DNA] = 60 μM ; [Hoechst] = 6 μM . Inset: plot [Ag] vs relative emission of Hoechst-bound DNA.

Determination of Ag uptake in the cell by HR-CS GFAAS

Table S2. Instrumental parameters and temperature program used in the determination of Ag in cell suspension by HR-CS GFAAS.

Wavelength (nm)	328.0683			
Read time (s)	5			
Integration mode of absorbance signal	Area			
Evaluation pixels	3 (CP $\pm$ 1)			
Purge gas	Argon			
Gas flow during atomization	Stop			
Working range (pg)	0 – 750			
Step	Name	Temp. ($^{\circ}\text{C}$)	Ramp. ($^{\circ}\text{C s}^{-1}$)	Hold (s)
1	Drying	125	7	30
2	Pyrolysis	800	60	35
3	Gas adaption	800	0	5
4	Atomize	1800	1500	5
5	Clean	2600	1000	3

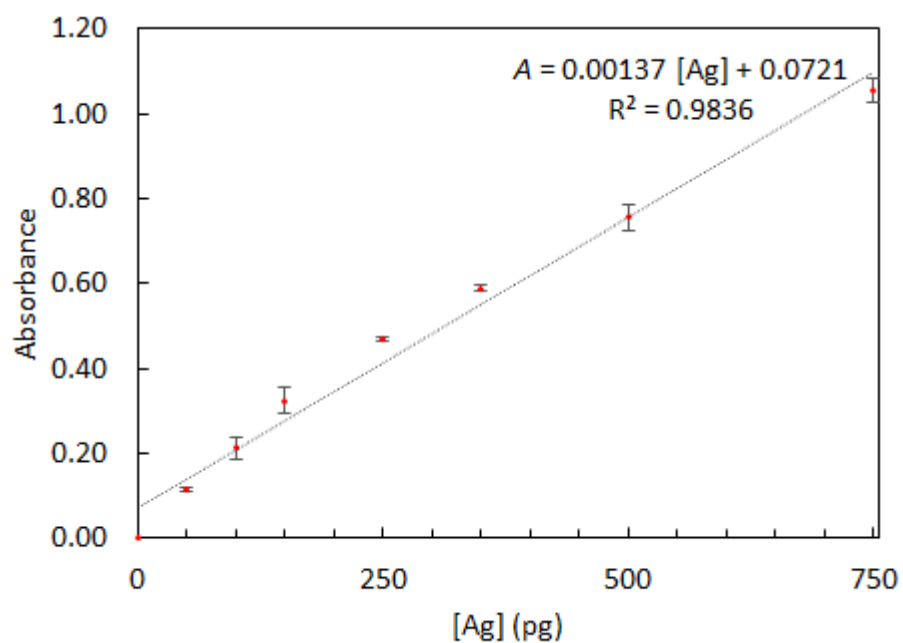


Fig. S18. Calibration curve obtained for Ag determination by HR-CS GFAAS.

Table S3. Percentage of Ag uptake in the cells determined by HR-CS GFAAS (n = 3).

Sample	[Ag] (pg)	Initial concentration (pg)	Uptake percentage (%)
1	2938 ± 172	9572 ± 119	30.7
	2916 ± 144		30.5
	3004 ± 92		31.4
2	2572 ± 170	7578 ± 211	33.9
	2594 ± 198		34.4
	2480 ± 135		32.7
3	3151 ± 105	8848 ± 408	35.6
	3138 ± 210		35.5
	3028 ± 291		34.2