

Electronic Supplementary Information

DNA and BSA Binding, Anticancer and Antimicrobial Properties of Co(II), Co(II/III), Cu(II) and Ag(I) Complexes of Arylhydrazones of Barbituric Acid

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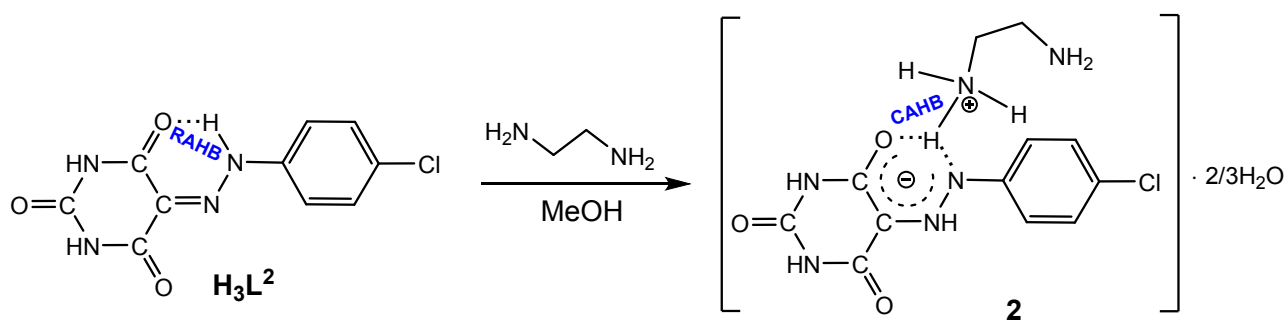
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1. Synthesis of 2



Scheme 1S. RAHB to CAHB transformations in the synthesis of **2**.

2. NMR and IR spectra of H_3L^2 , **1**, **2**, **7** and **8**.

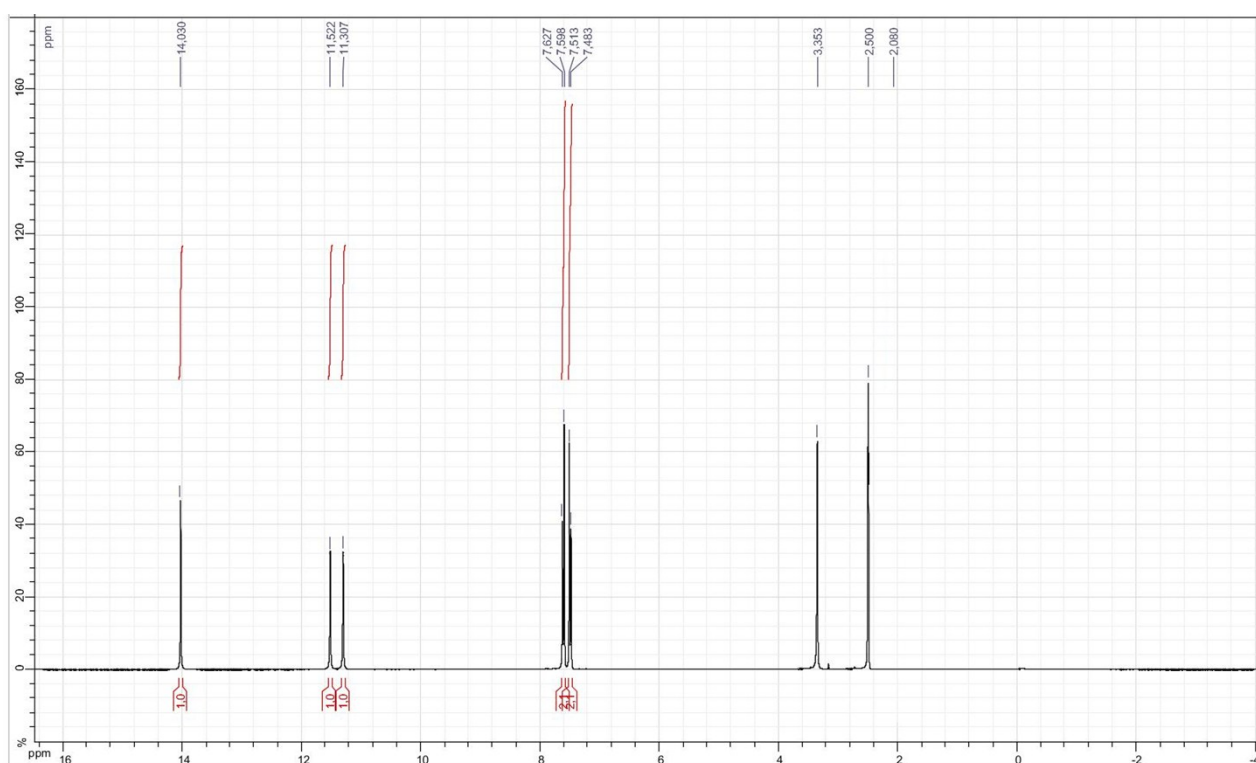


Figure 1S. ^1H NMR spectra of H_3L^2 .

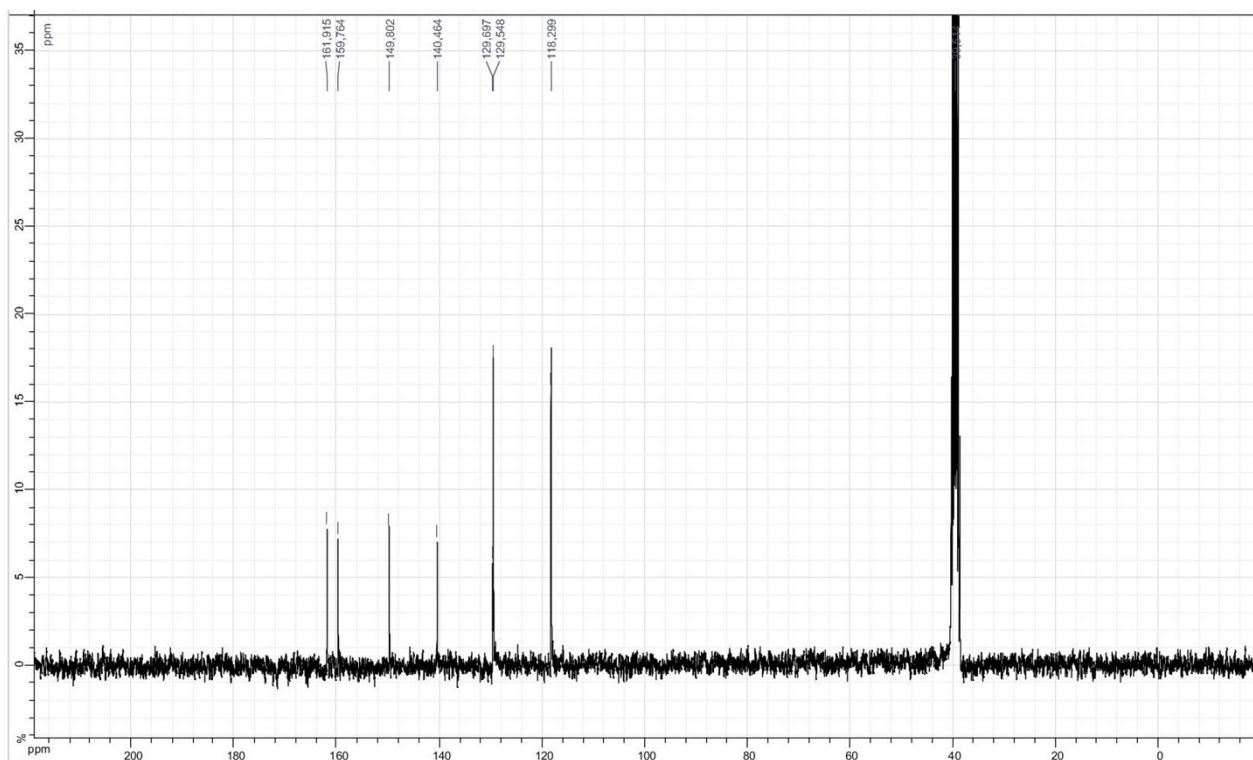


Figure 2S. ^{13}C NMR spectra of H_3L^2 .

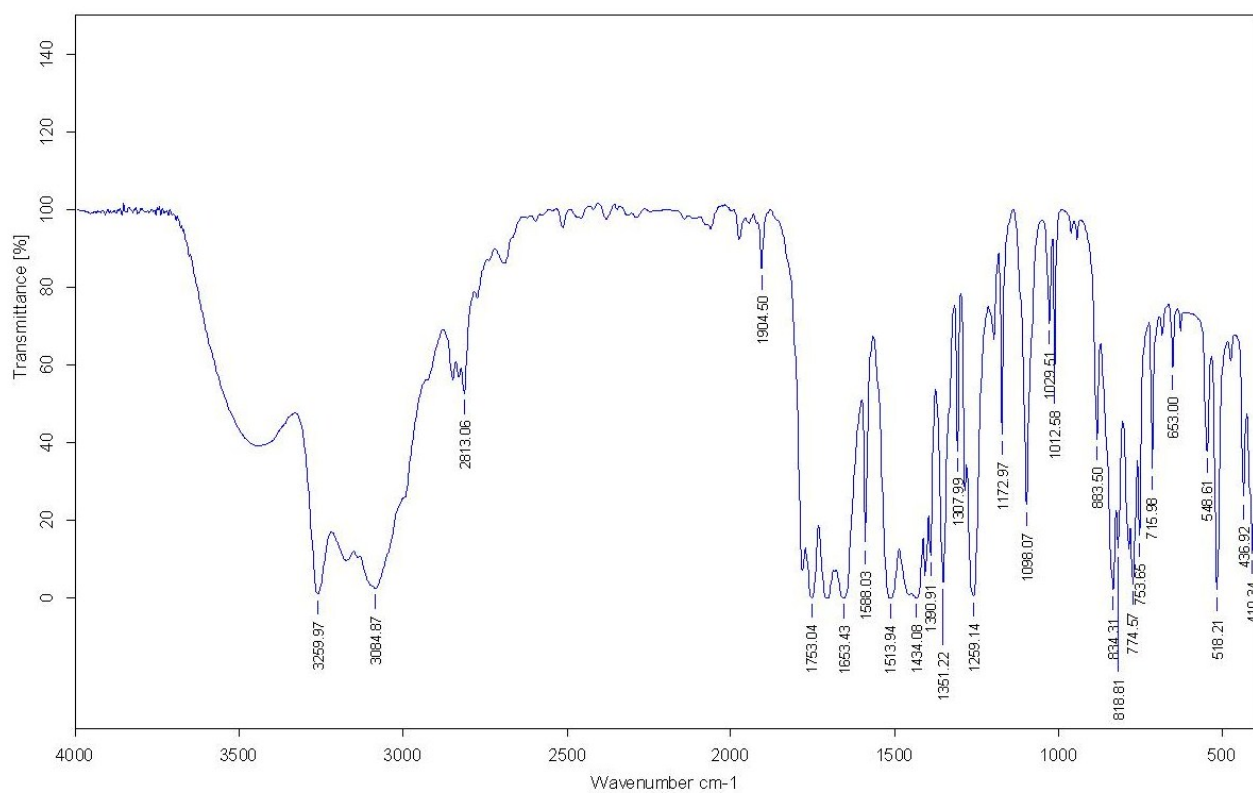


Figure 3S. IR spectra of H_3L^2 .

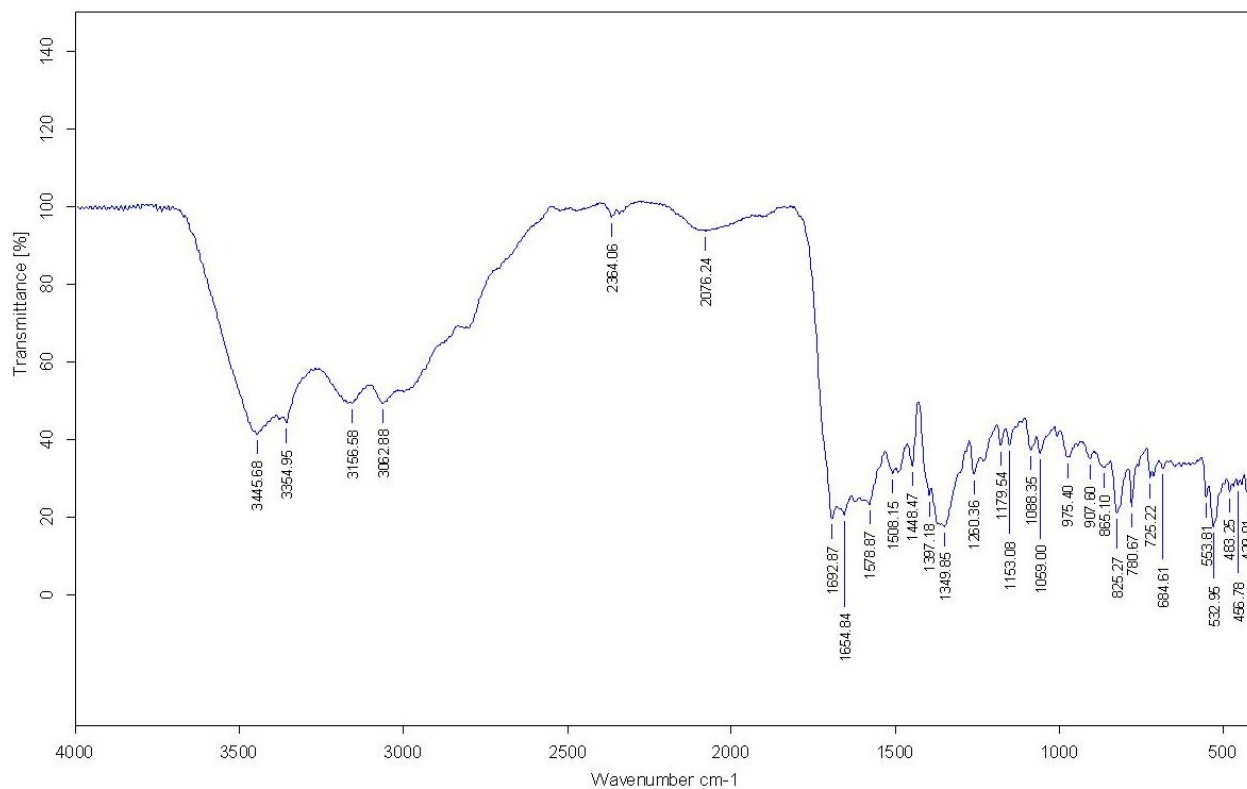


Figure 4S. IR spectra of 2.

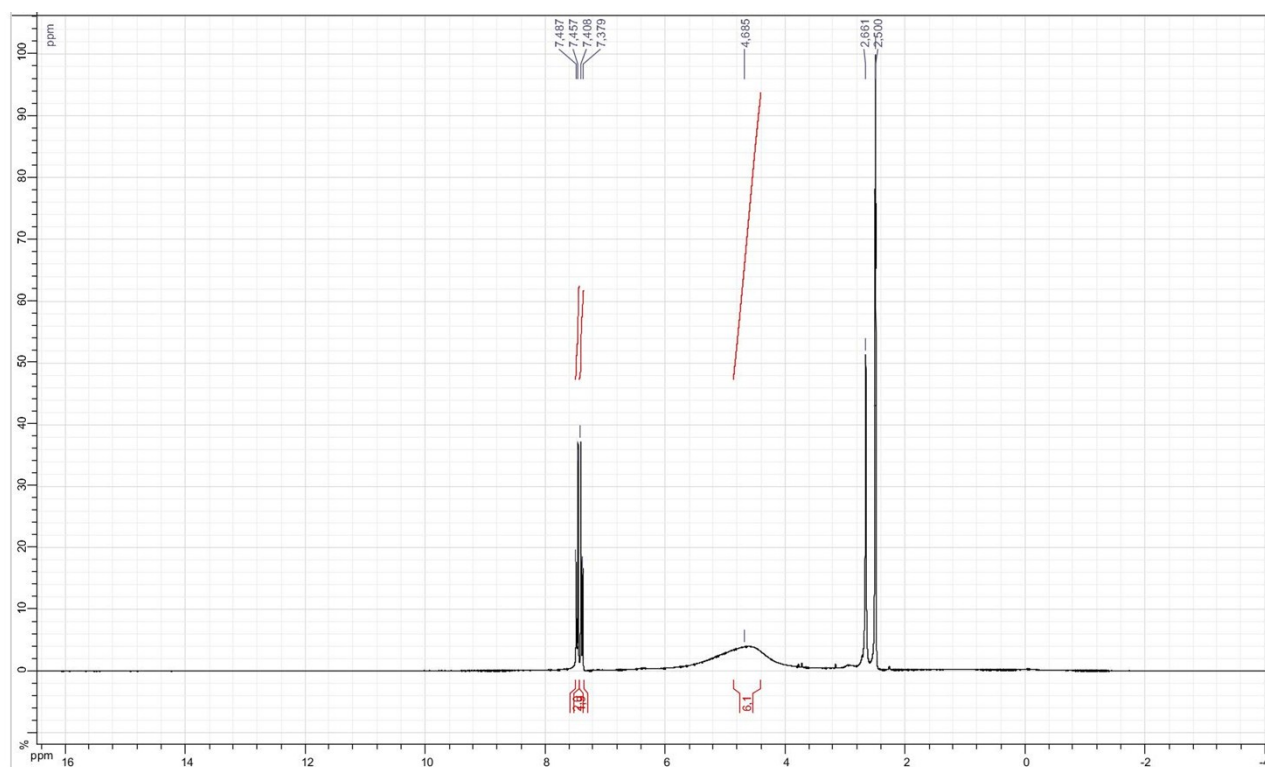


Figure 5S. ¹H NMR spectra of 2.

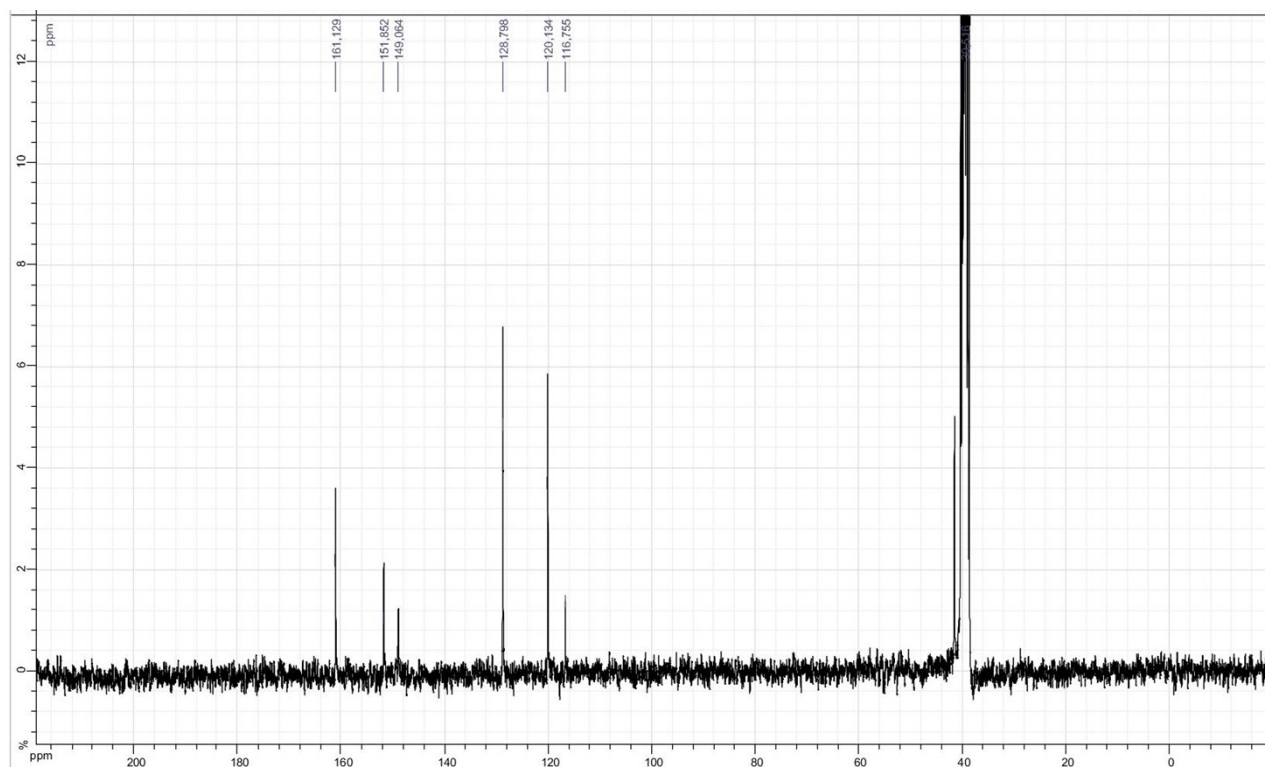


Figure 6S. ^{13}C NMR spectra of **2**.

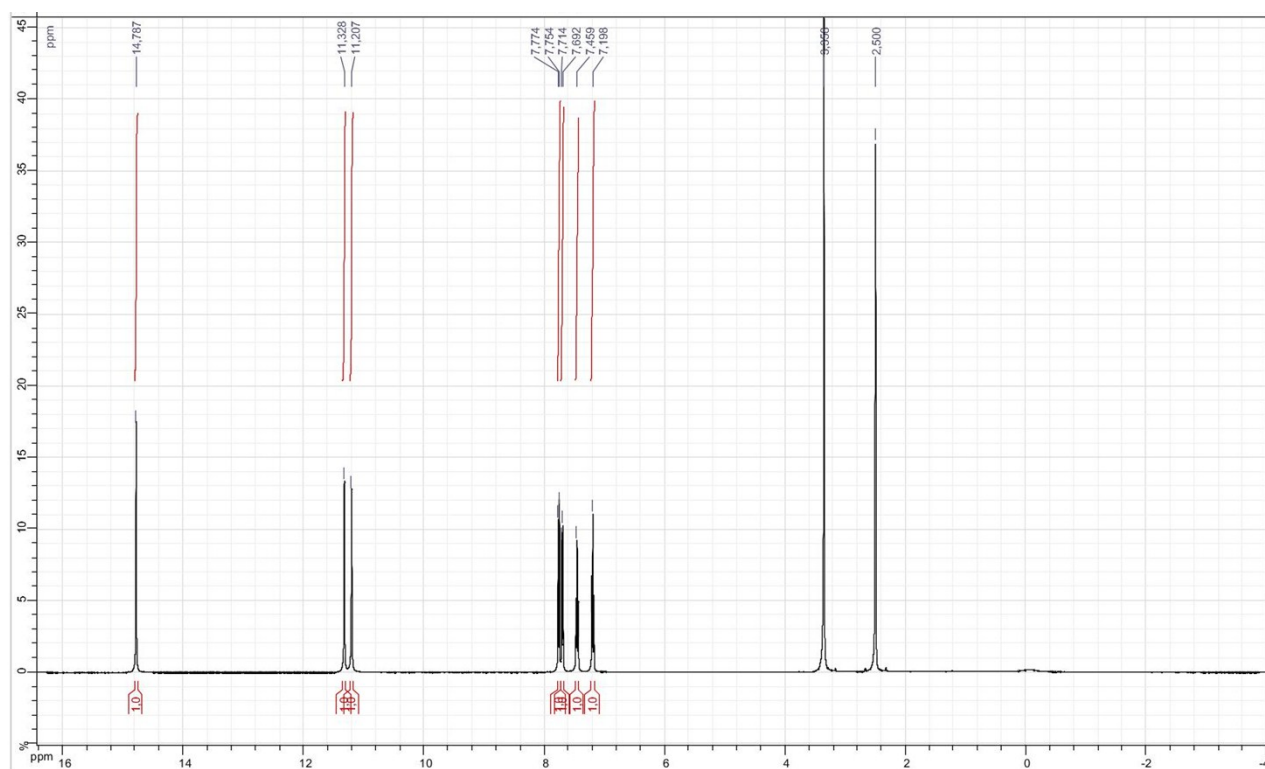


Figure 7S. ^1H NMR spectra of **7**.

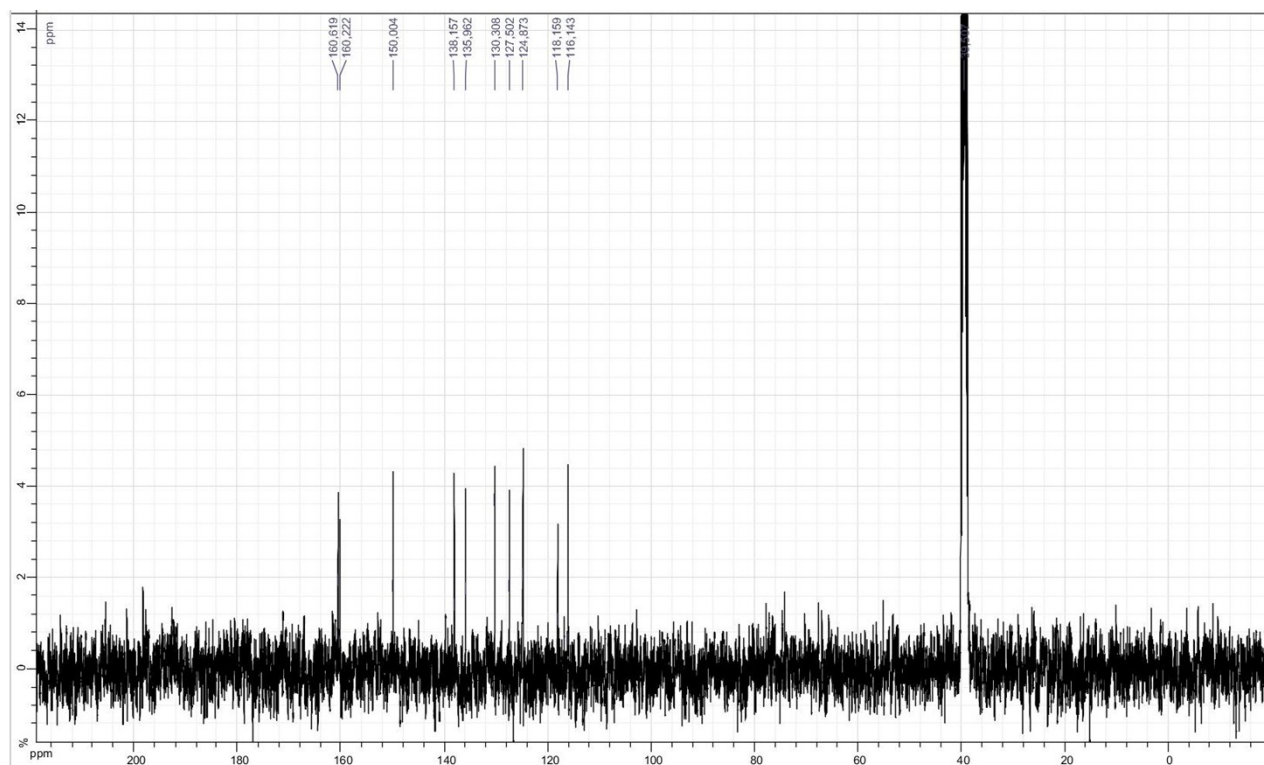


Figure 8S. ^{13}C NMR spectra of **7**.

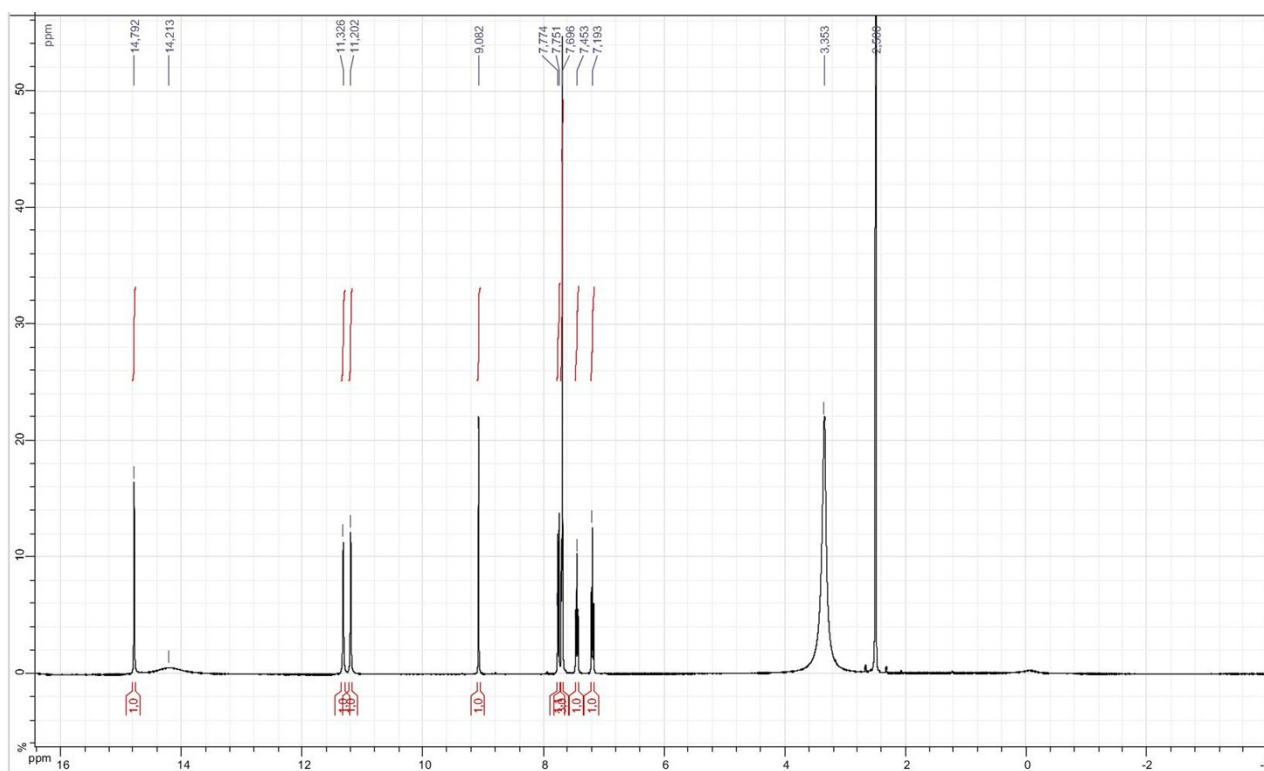


Figure 9S. ^1H NMR spectra of **8**.

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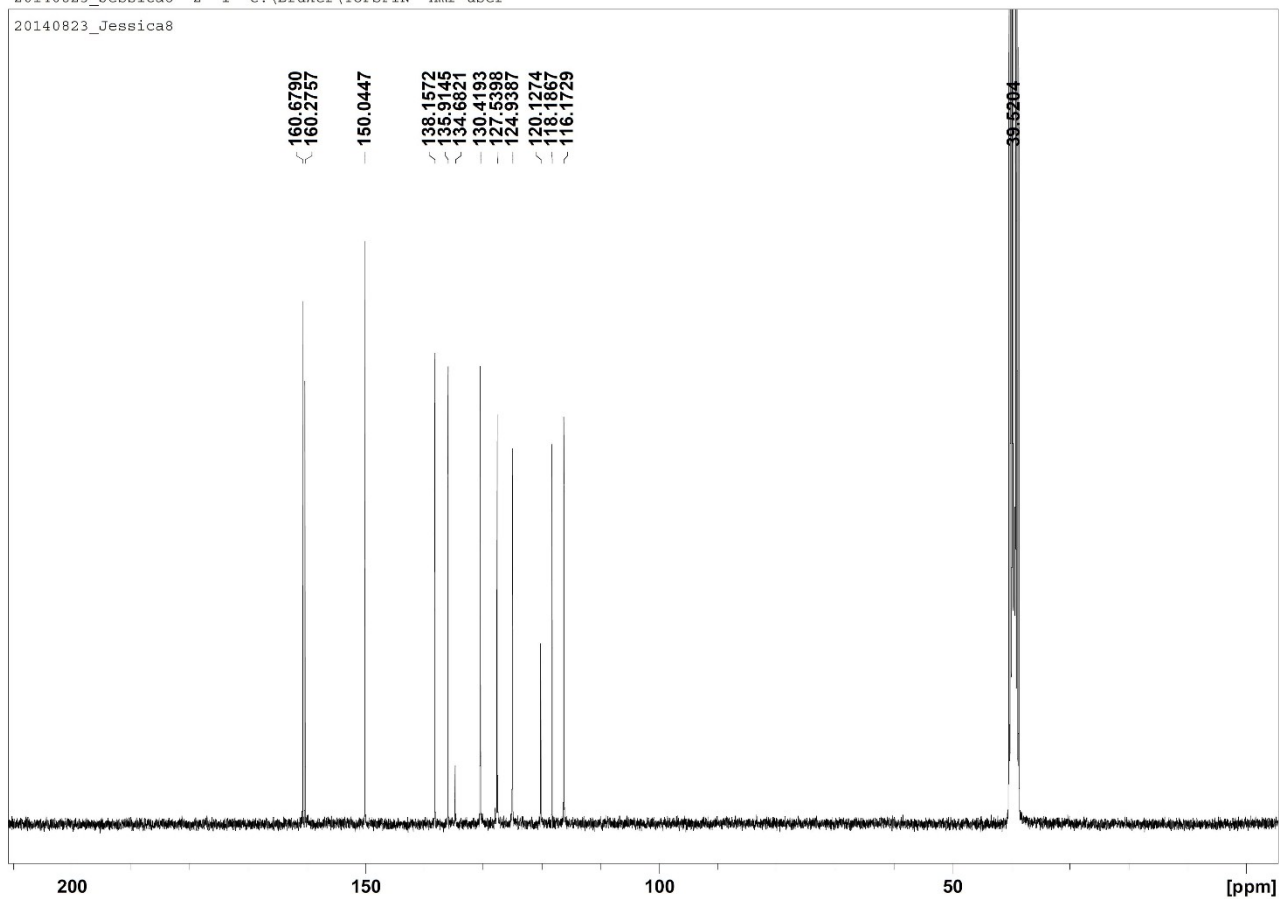


Figure 10S. ^{13}C NMR spectra of **8**.

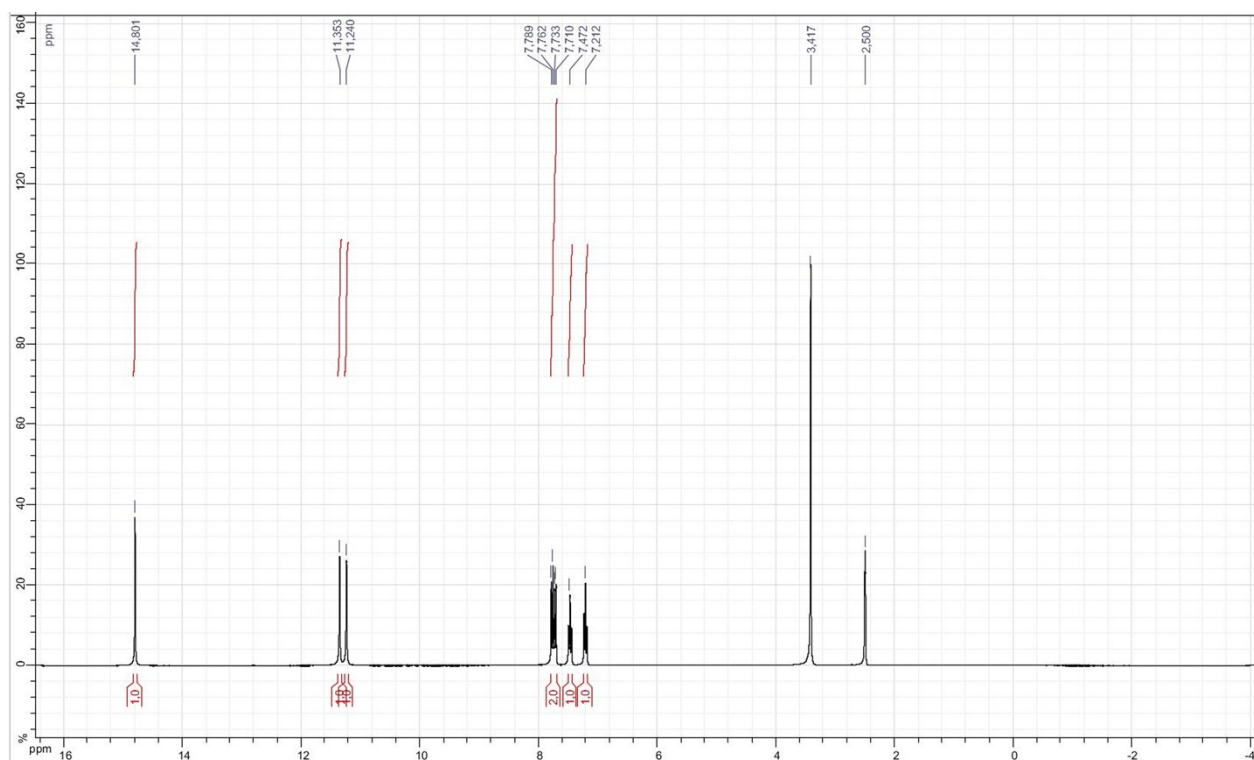


Figure 11S. ^1H NMR spectra of **1**.

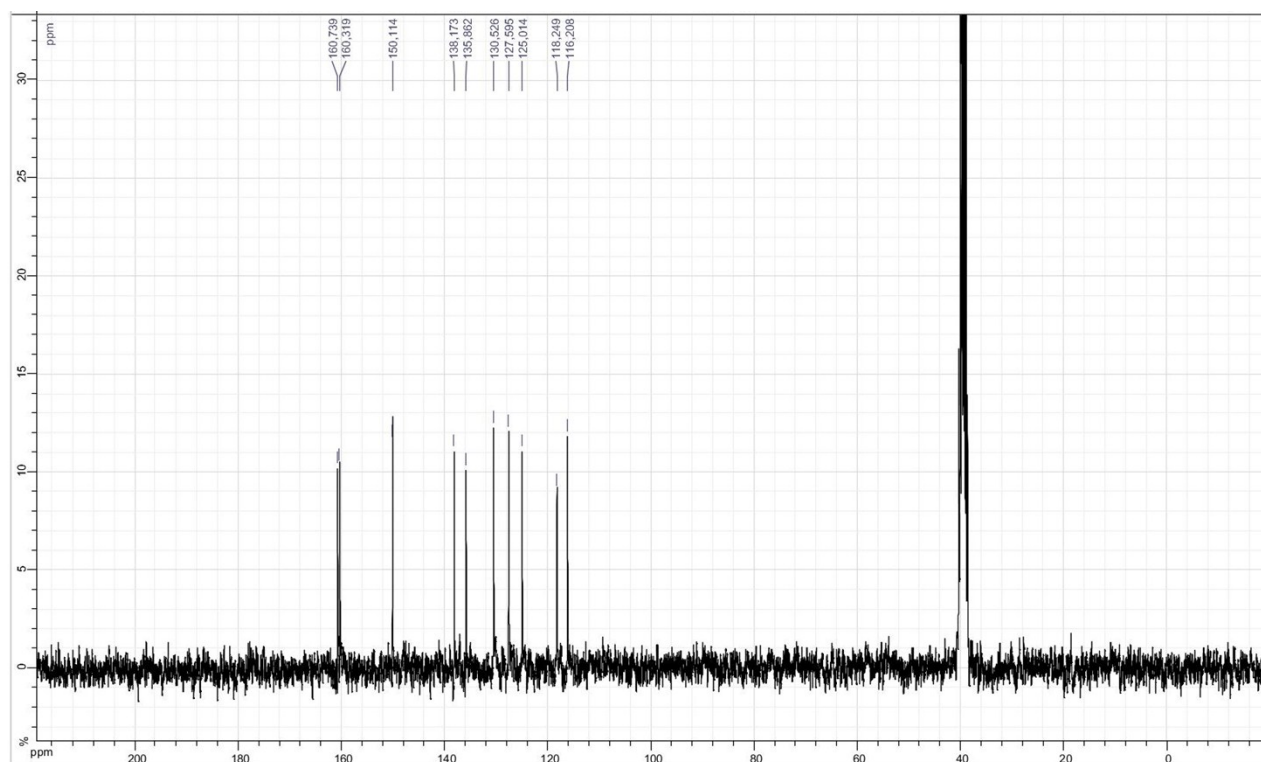


Figure 12S. ¹³C NMR spectra of 1.

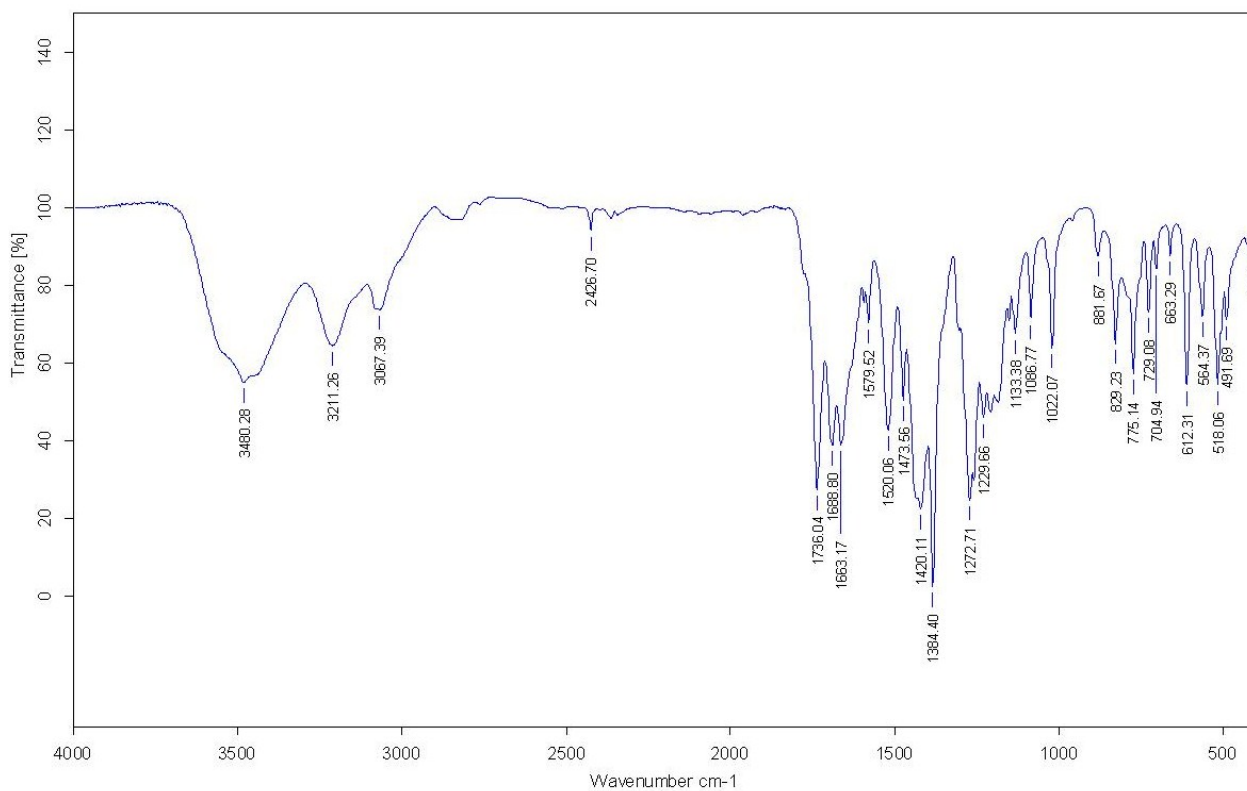


Figure 13S. IR spectra of 7.

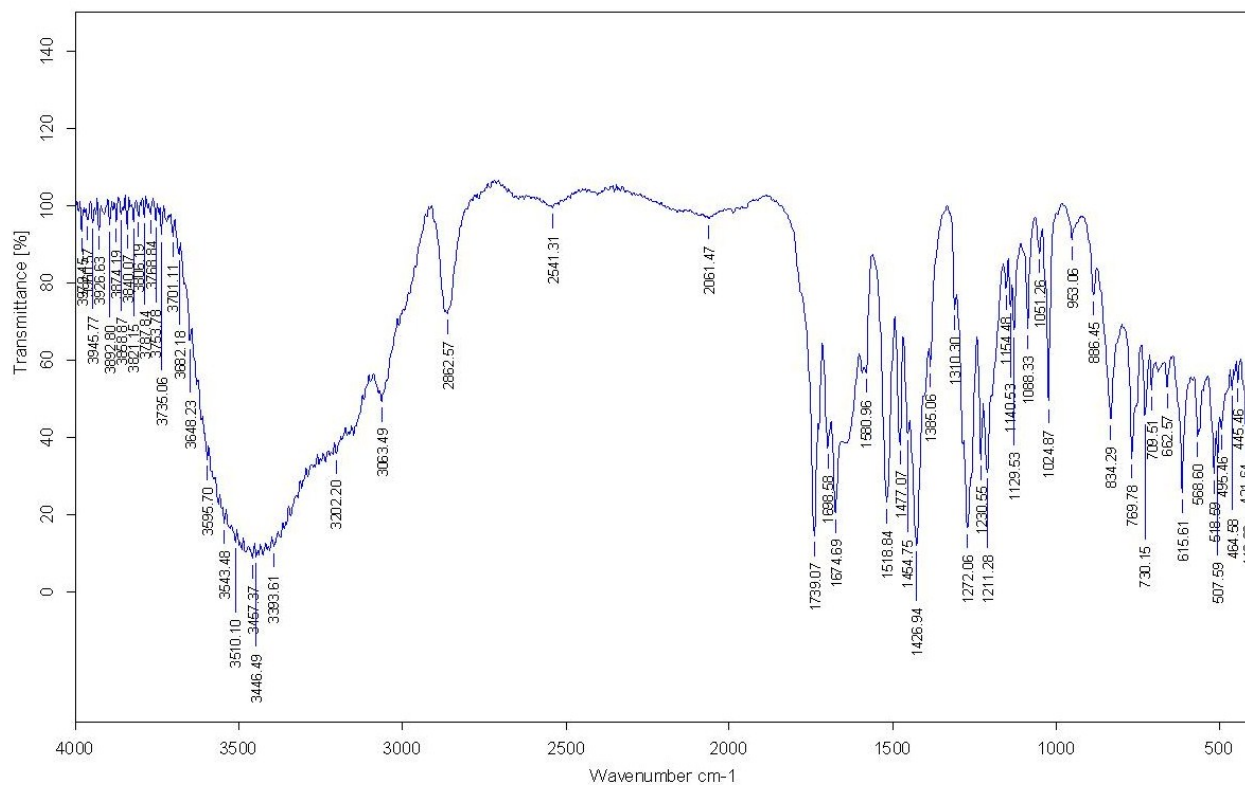


Figure 14S. IR spectra of 8.

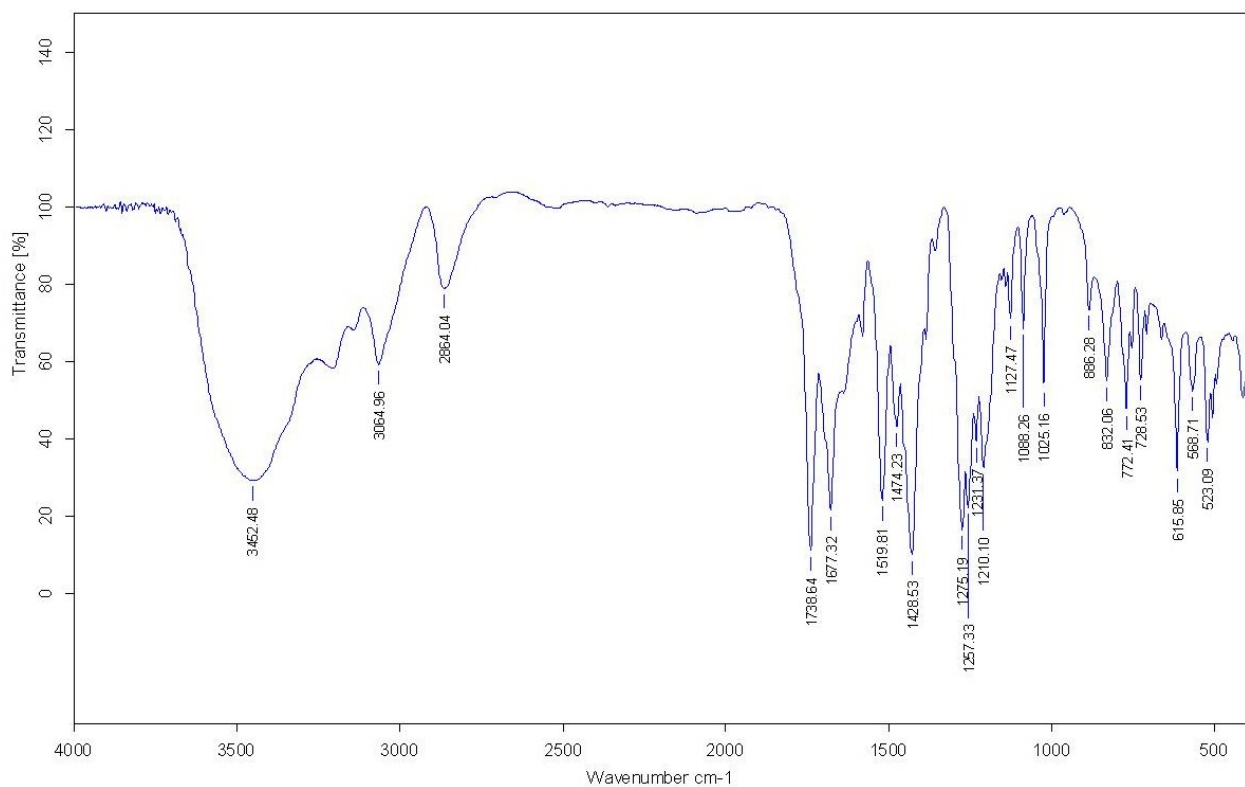


Figure 15S. IR spectra of 1.

3. X-ray analysis

Table 1S. Crystal data, experimental parameters and selected details of the refinement calculations of compounds **2**, **7** and **8**.

Compound	2	7	8
Empirical formula	C ₃₆ H ₄₉ Cl ₃ N ₁₈ O ₁₁	C ₁₀ H ₉ AgN ₄ O ₇ S	C ₁₃ H ₁₆ N ₆ O ₈ S
Formula weight	1016.28	437.14	416.38
Crystal system	Monoclinic	Triclinic	Triclinic
Space group	<i>P</i> 21/ <i>c</i>	<i>P</i> -1	<i>P</i> -1
<i>a</i> (Å)	10.4057(3)	8.4530(2)	7.1448(18)
<i>b</i> (Å)	21.2989(6)	9.1802(3)	11.280(4)
<i>c</i> (Å)	21.0249(6)	9.5557(3)	12.049(4)
α (deg)	90	69.666(3)	116.777(12)
β (deg)	100.811(1)	69.858(2)	98.803(11)
γ (deg)	90	79.279(2)	97.186(11)
<i>Z</i>	4	2	2
<i>V</i> (Å ³)	4577.0(2)	650.93(4)	835.8(4)
<i>T</i> (K)	296	150	150
ρ_{calc} (g/cm ³)	1.475	2.230	1.655
μ (Mo K α) (mm ⁻¹)	0.279	1.757	0.256
Rfl collected/unique/obs	66617/5164/3794	11649/4749/4228	13696/3413/2364
<i>R</i> ^{1a} (<i>I</i> ≥ 2 σ)	0.0857	0.0465	0.0590
w <i>R</i> ^{2b} (<i>I</i> ≥ 2 σ)	0.2558	0.1401	0.1523
GOF on <i>F</i> ²	1.135	1.093	0.957

^[a] $R1 = \sum ||Fo| - |Fc|| / \sum |Fo|$. ^[b] $wR2 = [\sum [w(Fo^2 - Fc^2)^2] / \sum [w(Fo^2)^2]]^{1/2}$.

Table 2S. Selected distances (Å) and angles (°) for compounds **2**, **7** and **8**.

2		7		8	
C8–O1	1.232(7)	C7–N2	1.327(4)	C7–O1	1.232(4)
C9–O2	1.238(7)	C8–O1	1.239(3)	C9–O2	1.217(4)
C7–N2	1.395(8)	C9–O2	1.224(3)	C8–N2	1.326(4)
N1–N2	1.202(8)	O1–Ag1	2.589(3)	C10–O3	1.231(4)
C4–N1	1.498(9)	O3–Ag1	2.566(2)	N1–N2	1.308(4)
C10–O3	1.229(7)	O10–Ag1	2.318(3)	N1–N2–C8	120.5(3)
N2–N1–C4	111.5(7)	O11–Ag1	2.343(2)	N2–C8–C7	124.2(3)
N1–N2–C7	118.1(7)	O13–Ag1	2.359(2)	O1–C7–C8	123.8(3)

Table 3S. Hydrogen bond interactions (Å, °) in complexes **2**, **7** and **8**.

D–H···A	<i>d</i> (H···A)	<i>d</i> (D···A)	∠(D–H···A)	Symmetry operation
2				
O1W···H1A···O9	1.90(4)	2.758(7)	156(6)	<i>x, 1/2-y, 1/2+z</i>
O2W···H2A···O6	1.91(7)	2.801(6)	167(7)	<i>x, 1/2-y, -1/2+z</i>
O2W···H2B···O1	2.12(8)	2.821(6)	134(8)	<i>intra</i>
O2W···H2B···N1	2.21(6)	2.972(10)	143(9)	<i>intra</i>
N3···H3N···O5	1.90(11)	2.900(6)	165(8)	<i>x, 1/2-y, -1/2+z</i>
N4···H4N···O6	2.08(4)	2.954(6)	170(11)	<i>x, 1/2-y, -1/2+z</i>
N7···H7N···O3	1.98(3)	2.872(6)	177(15)	<i>l+x, 1/2-y, 1/2+z</i>
N8···H8N···O1	1.84(11)	2.814(6)	174(10)	<i>x, 1/2-y, 1/2+z</i>
N11···H11N···O7	1.96(12)	2.824(6)	166(8)	<i>l-x, l-y, l-z</i>
N12···H12N···O9	2.08(12)	2.938(6)	164(9)	<i>-x, l-y, l-z</i>
N31···H31C···O2	2.08(7)	2.868(7)	148(9)	<i>intra</i>
N31···H31D···O1W	2.19(11)	3.042(9)	148(10)	<i>intra</i>
N32···H32C···O2	2.52(11)	3.118(7)	122(8)	<i>intra</i>
N32···H32C···N2	2.38(11)	3.296(8)	164(9)	<i>intra</i>
N32···H32D···O5	2.28(10)	3.011(7)	133(8)	<i>intra</i>
N32···H32D···N6	2.31(11)	3.193(7)	154(9)	<i>intra</i>
N32···H32E···N35	1.88(11)	2.836(8)	172(5)	<i>-l+x,y,z</i>
N33···H33C···O5	2.34	3.175(8)	156	<i>intra</i>
N33···H33D···O8	2.37	3.094(7)	138	<i>intra</i>
N34···H34C···N31	1.84(11)	2.811(9)	168(10)	<i>-x, l/2+y, 3/2-z</i>
N34···H34D···O2W	1.90(8)	2.785(8)	170(8)	<i>l-x, l/2+y, 3/2-z</i>
N34···H34E···O8	2.38(11)	2.940(7)	125(9)	<i>intra</i>
N34···H34E···N9	2.28(11)	3.074(8)	159(9)	<i>intra</i>
N35···H35C···O4	2.42(12)	2.992(7)	124(10)	<i>intra</i>
N36···H36C···O1W	2.40(11)	2.991(9)	122(7)	<i>l-x, l/2+y, 3/2-z</i>
N36···H36D···O7	1.93(8)	2.810(7)	170(8)	<i>intra</i>
N36···H36E···O4	2.13(10)	2.770(7)	124(8)	<i>intra</i>
N36···H36E···N5	2.13(12)	2.976(8)	149(10)	<i>intra</i>
7				
N1···H1N···O1	1.87(4)	2.613(4)	142(6)	<i>intra</i>
N1···H1N···O13	2.22(6)	2.815(4)	125(5)	<i>intra</i>
N3···H3N···O12	2.17(3)	3.037(4)	166(6)	<i>l-x, l-y, l-z</i>
N4···H4N···O13	2.50(6)	3.059(4)	121(5)	<i>x, l+y, z</i>
N4···H4N···O2	2.09(6)	2.912(4)	153(5)	<i>l-x, 2-y, 2-z</i>
O10···H10A···O11	1.98(5)	2.911(5)	159(5)	<i>-x, l-y, l-z</i>
O10···H10B···O2	1.86(5)	2.815(4)	169(4)	<i>x,y, -l+z</i>
8				
O1W···H1W1···O12	2.02(4)	2.830(4)	159(4)	<i>intra</i>
N1···H1N···O1	2.01(5)	2.650(5)	124(4)	<i>intra</i>
N1···H1N···O13	1.94(4)	2.714(4)	138(4)	<i>intra</i>
O1W···H1W2···O11	2.27(4)	2.934(4)	146(5)	<i>l+x,y,z</i>
O2W···H2W1···O3	1.88(3)	2.842(3)	166(4)	<i>l-x, l-y, -z</i>
N3···H3N···O1W	1.87(4)	2.776(4)	178(5)	<i>-l+x,y, -l+z</i>
O2W···H2W2···O11	1.83(2)	2.718(4)	175(3)	<i>l-x, l-y, l-z</i>
N4···H4N···O3	2.09(4)	2.911(4)	175(5)	<i>l-x, l-y, -z</i>
N11···H11N···O2W	1.84(4)	2.706(4)	172(4)	<i>intra</i>
N12···H12N···O13	2.21(4)	2.798(4)	127(4)	<i>-l+x,y,z</i>
N12···H12N···O2	2.19(5)	2.888(5)	140(3)	<i>-x,-y,-z</i>

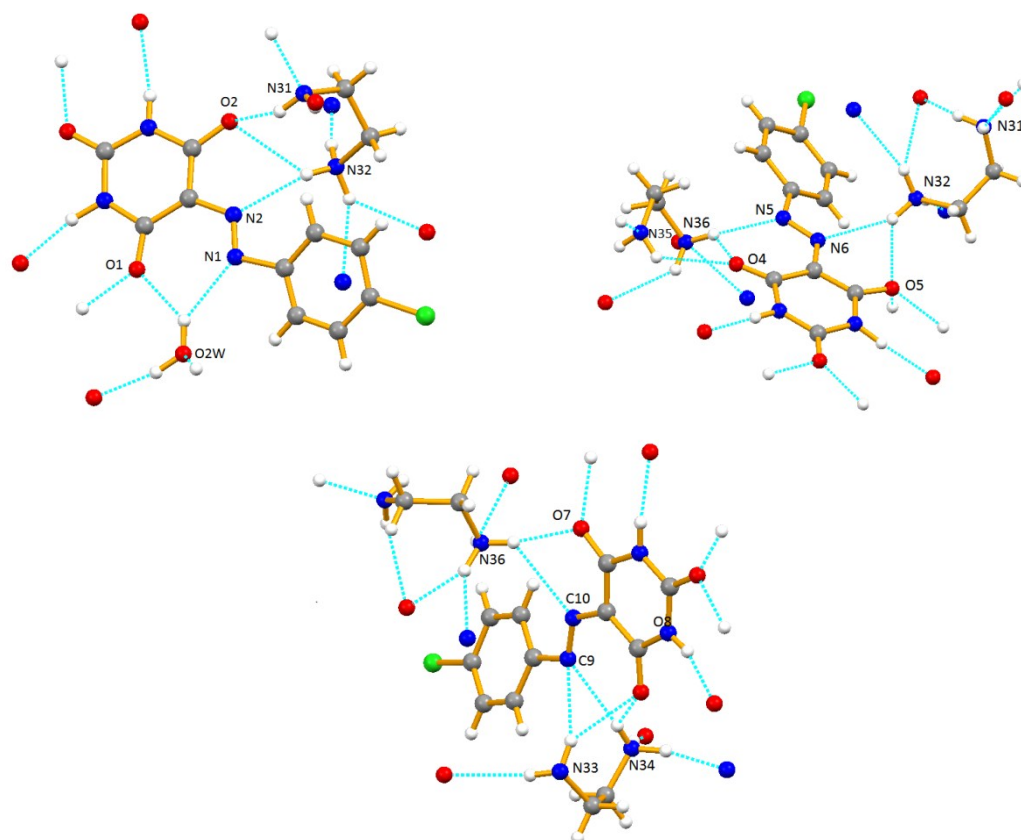
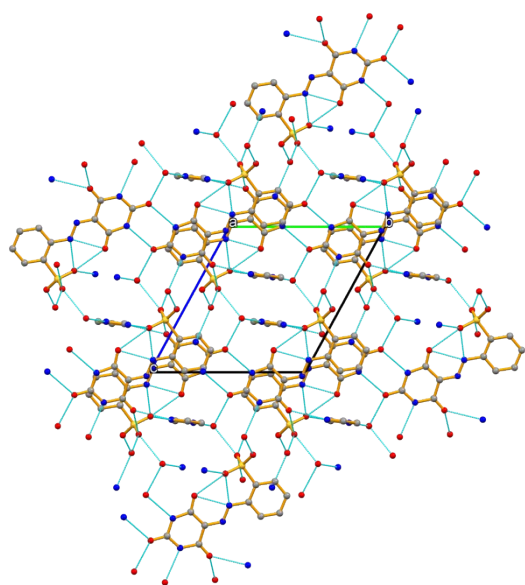
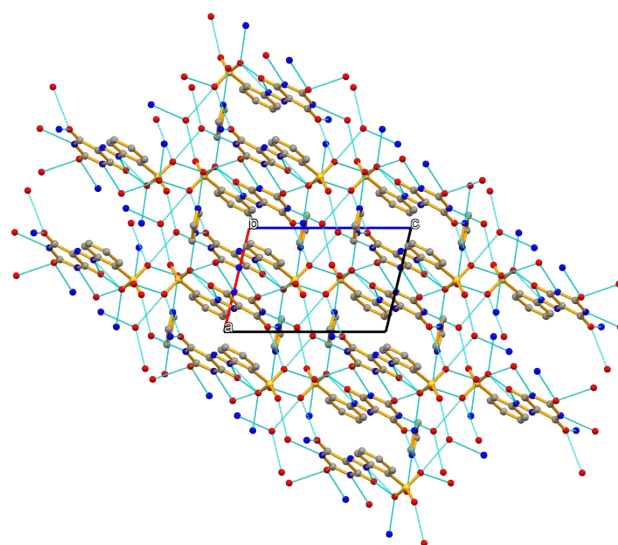


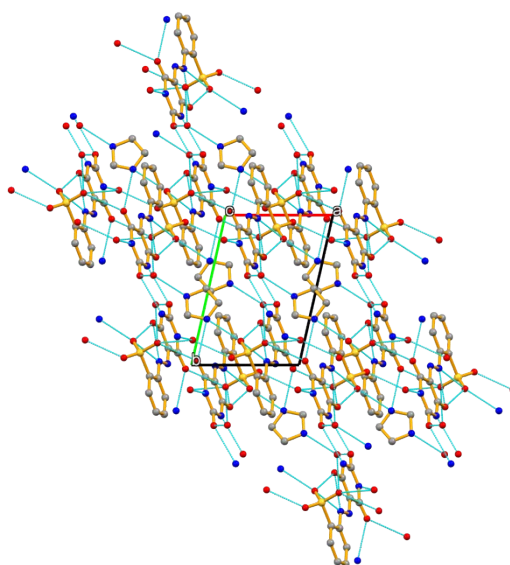
Figure 16S. Hydrogen bond interactions in **2** (in dashed blue lines; see also Table 3S) grouped by hydrazone anion and monoprotonated ethylenediamine cation.



(a)



(b)



(c)

Figure 17S. 3D network arrangement in **2** viewed down the crystallographic *a*, *b* and *c* axis, respectively. Hydrogen atoms were omitted for clarity.

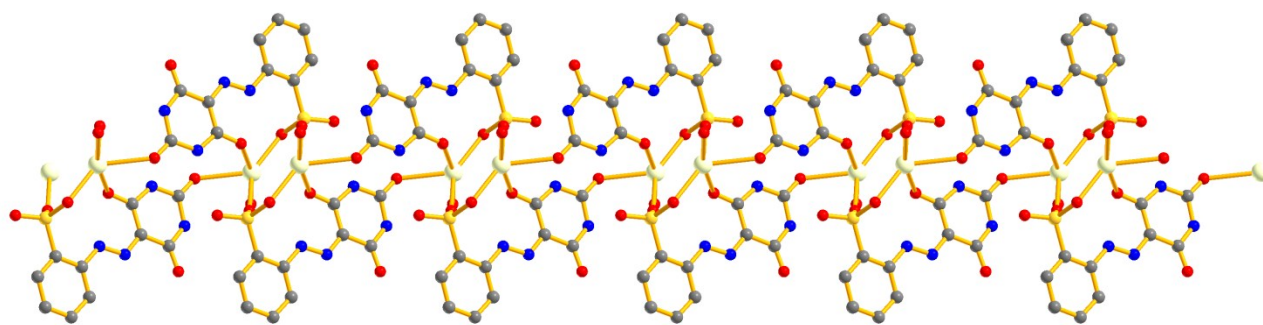


Figure 18S. 1D polymeric chain arrangement in **7**. Hydrogen atoms were omitted for clarity.

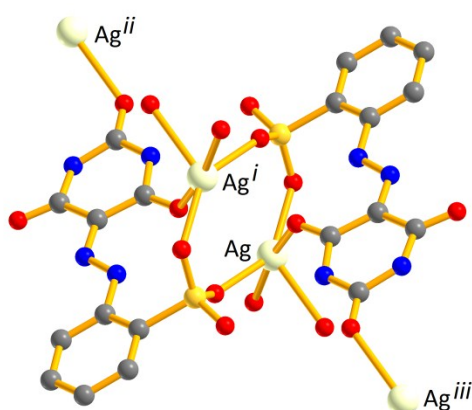


Figure 19S. The bridging mode of the sulfonyl groups in **7** which generate $\{Ag_2S_2O_4\}$ dimetallic cores. i) 1-x,1-y,1-z; ii) x,-1+y,z; iii) 1-x,2-y,1-z

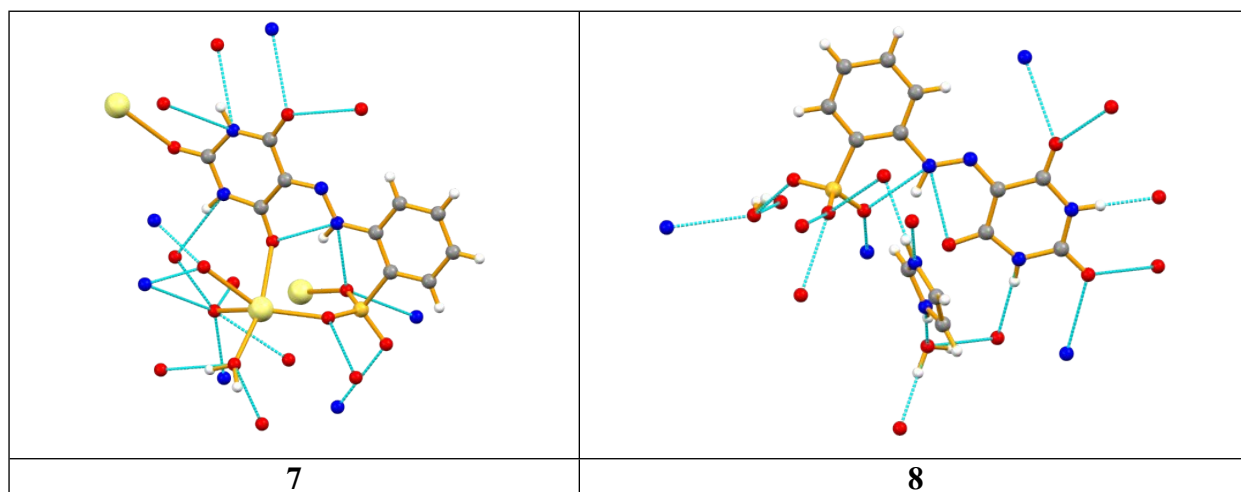


Figure 20S. Hydrogen bond interactions (in dashed blue lines; see also Table 3S) in **7** and **8**.

4. Electronic absorption spectra

Table 4S. Electronic absorption spectral data of proligands **H₄L¹** and **H₃L²** and complexes **1–9**.

Compound	λ_{max} (nm)	ϵ (M ⁻¹ cm ⁻¹) $\times 10^{-3}$
H₄L¹	414	37.6
	266	8.10
H₃L²	388	37.5
	262	7.51
1	389	50.1
	234	2.41
2	388	76.5
	262	16.9
3	480	23.0
	366	12.5
	269	15.7
4	424	72.2
	350	25.3
	269	63.4
5	393	18.3
	350	14.9
	260	16.3
6	388	21.6
	265	6.04
7	388	21.1
8	388	33.3
	262	13.2
9	384	64.1
	262	20.2

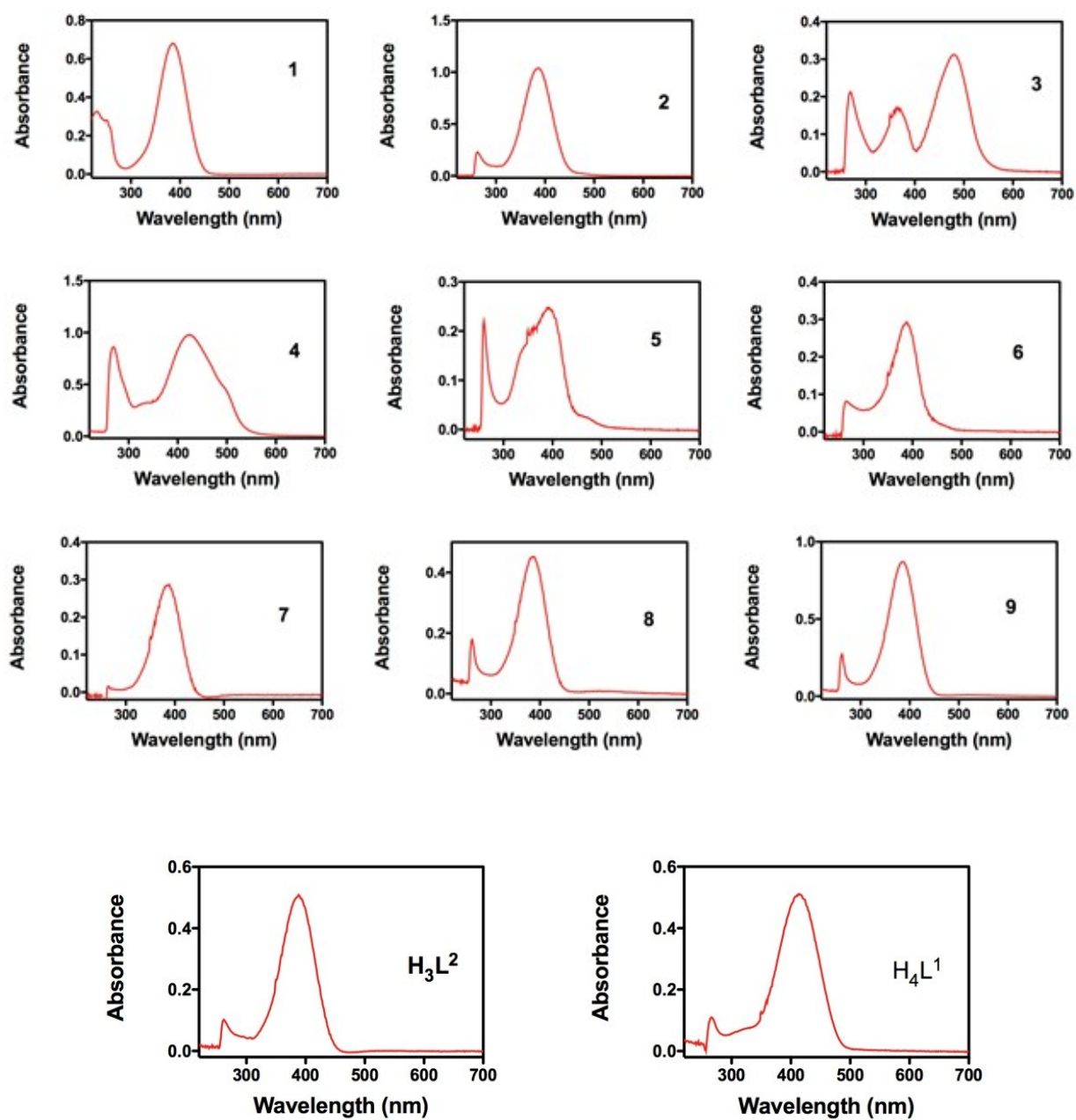
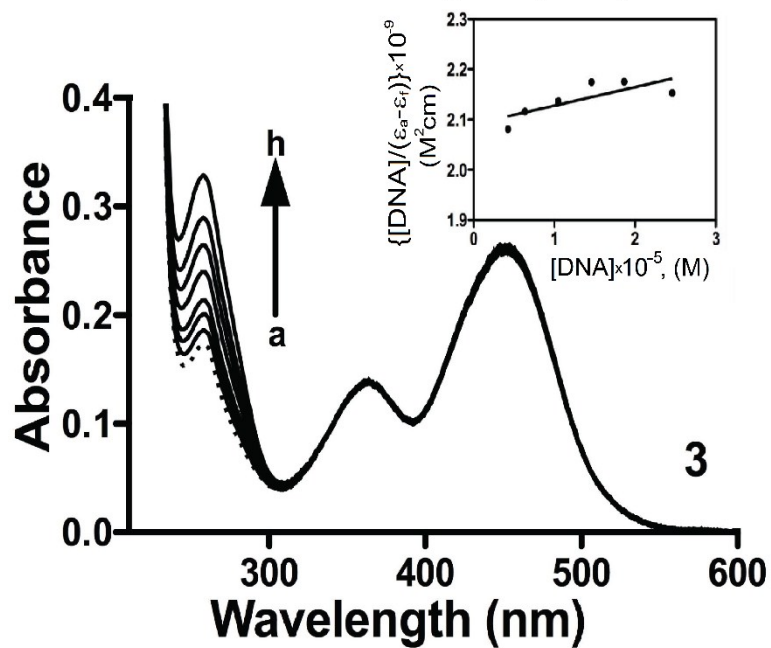
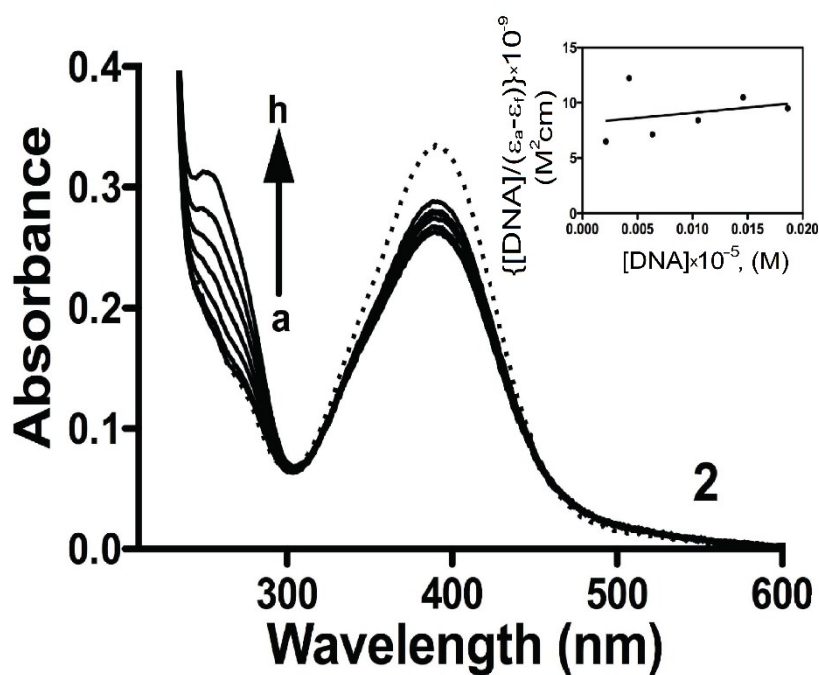
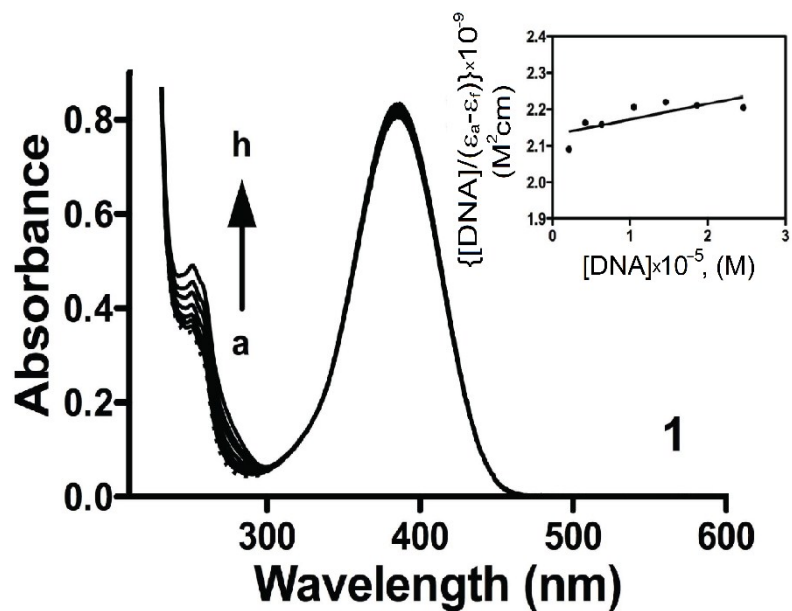
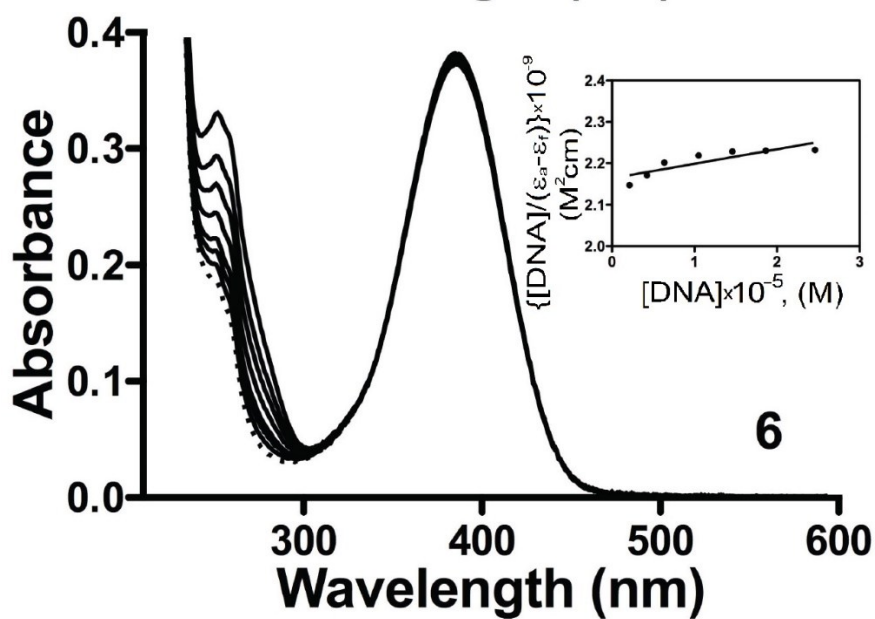
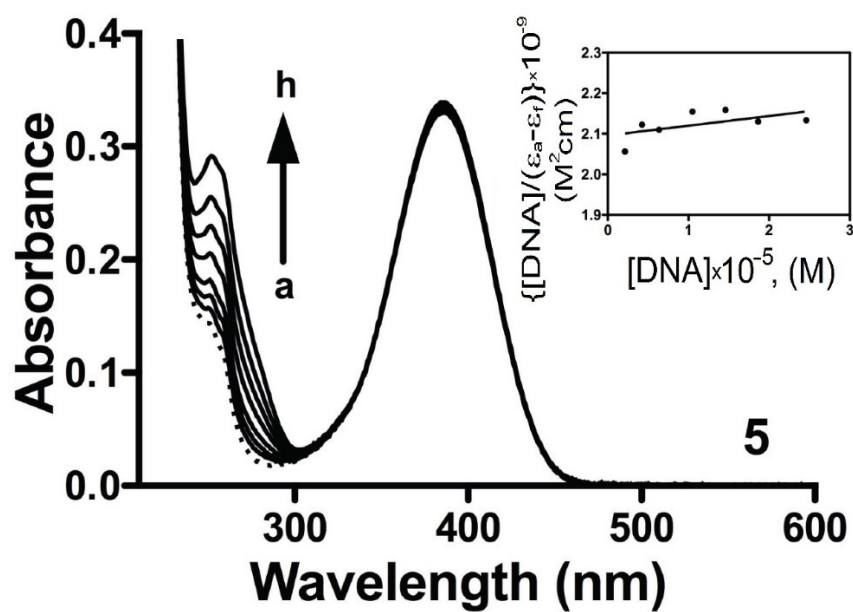
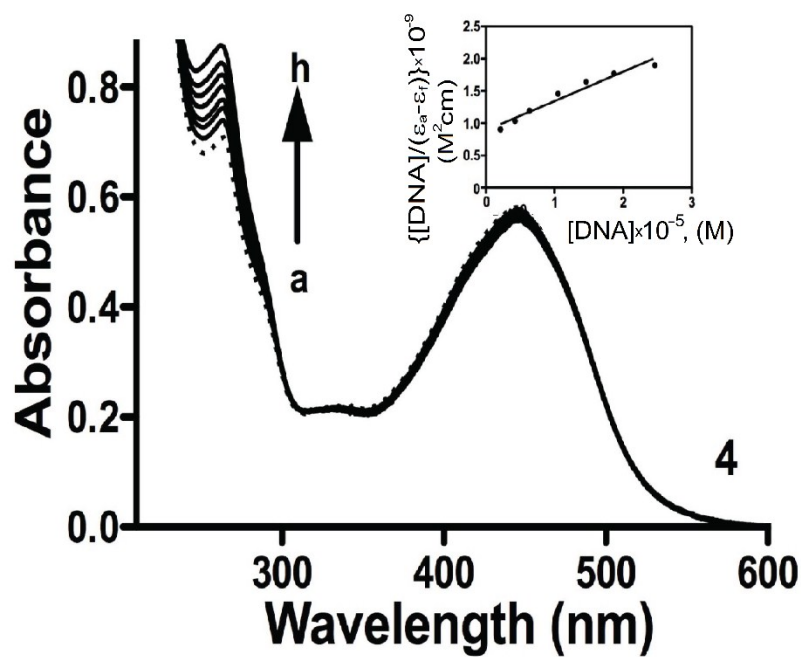
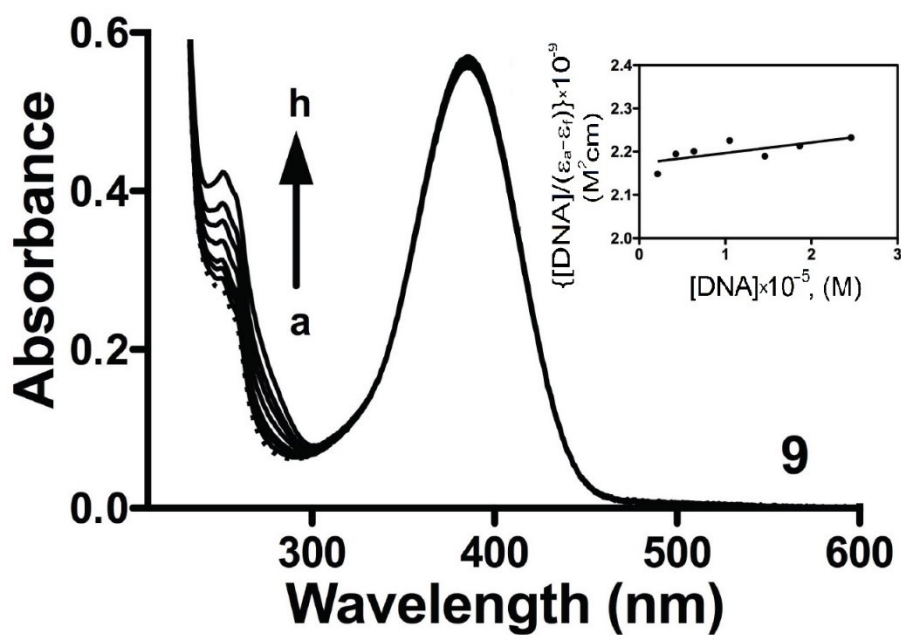
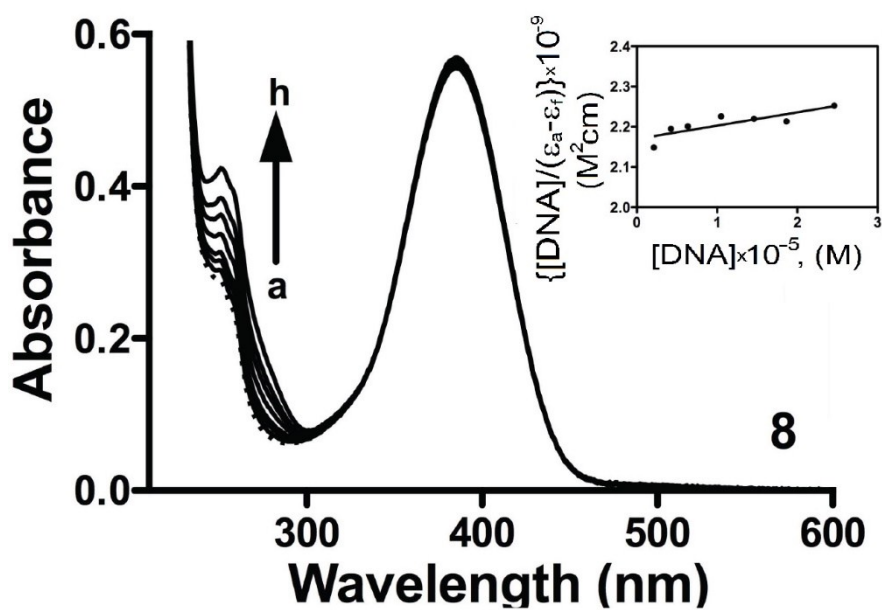
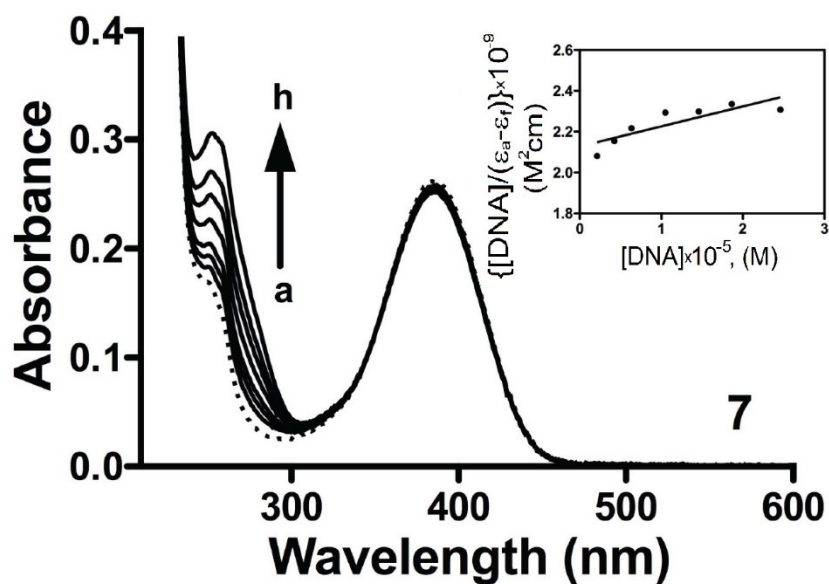


Figure 21S. UV-vis spectra for different compounds 15 μ M in DMSO (**H₄L¹**, **H₃L²**, **2-9**) and water (**1**) solution.







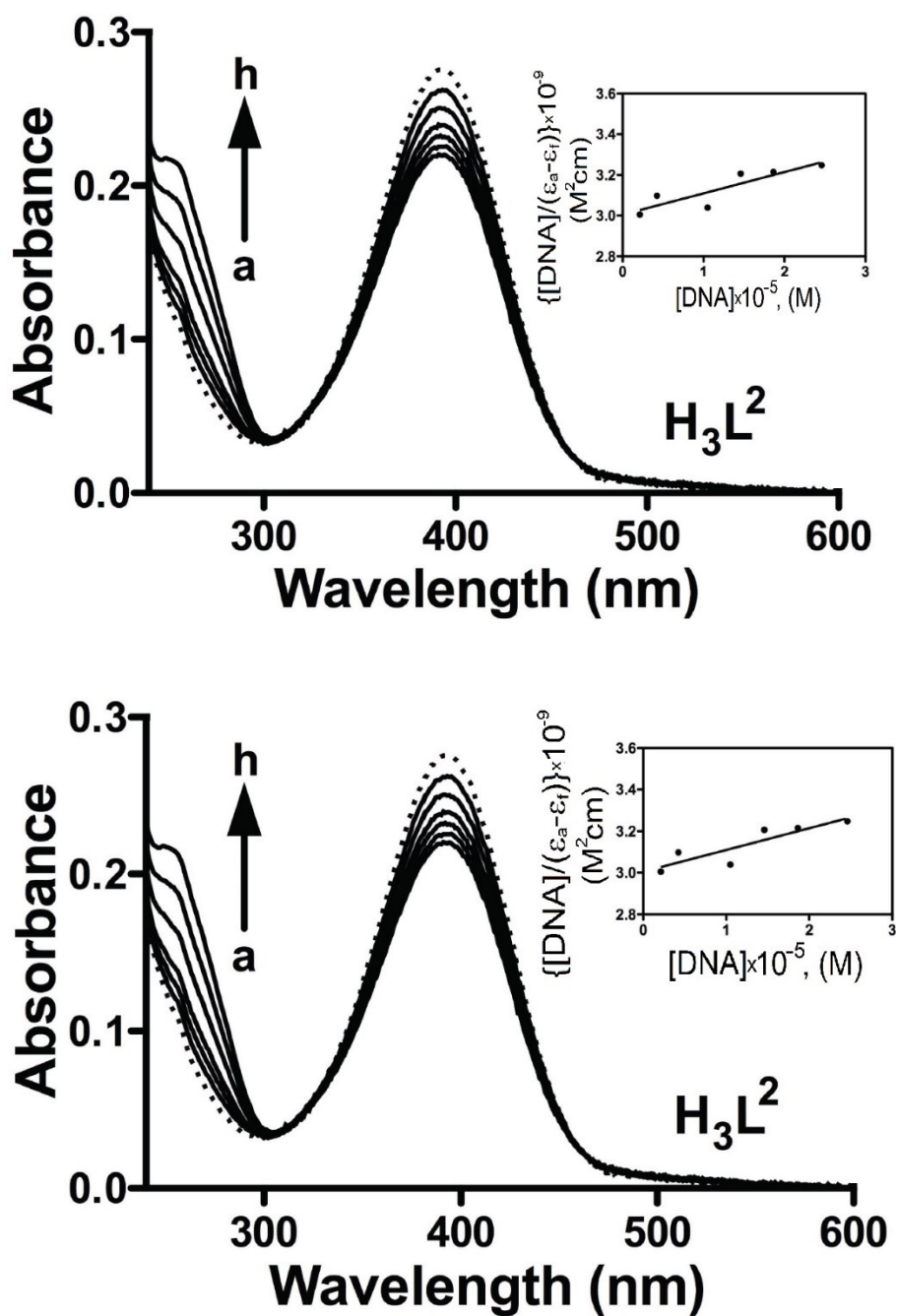
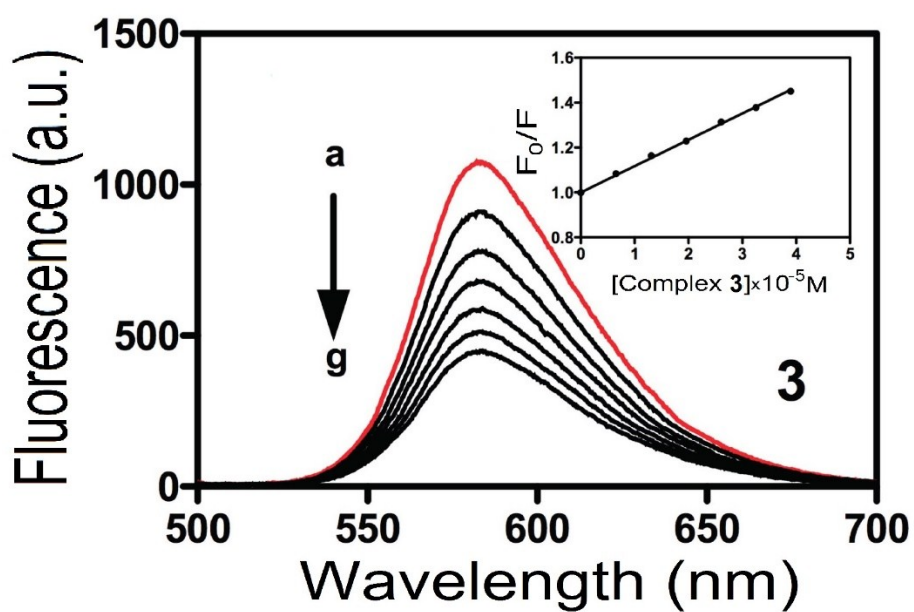
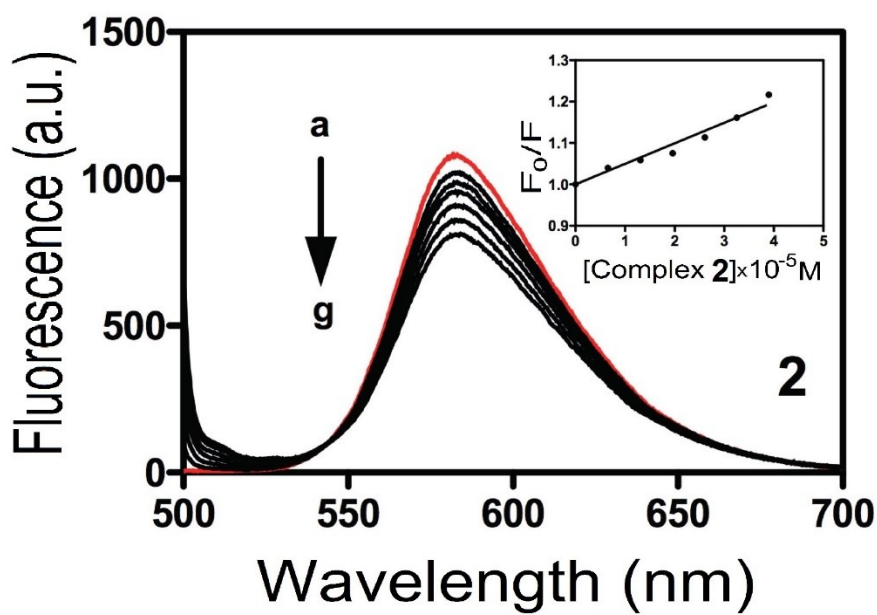
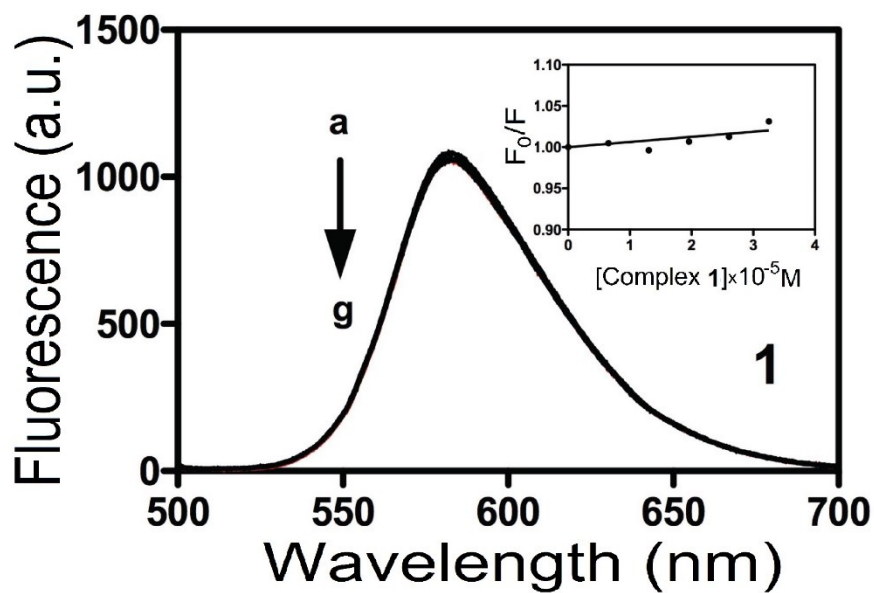
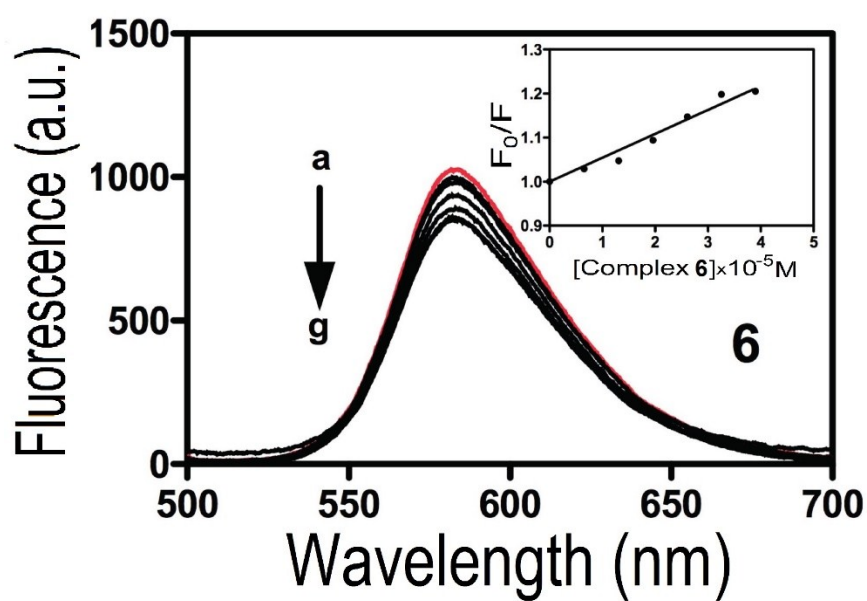
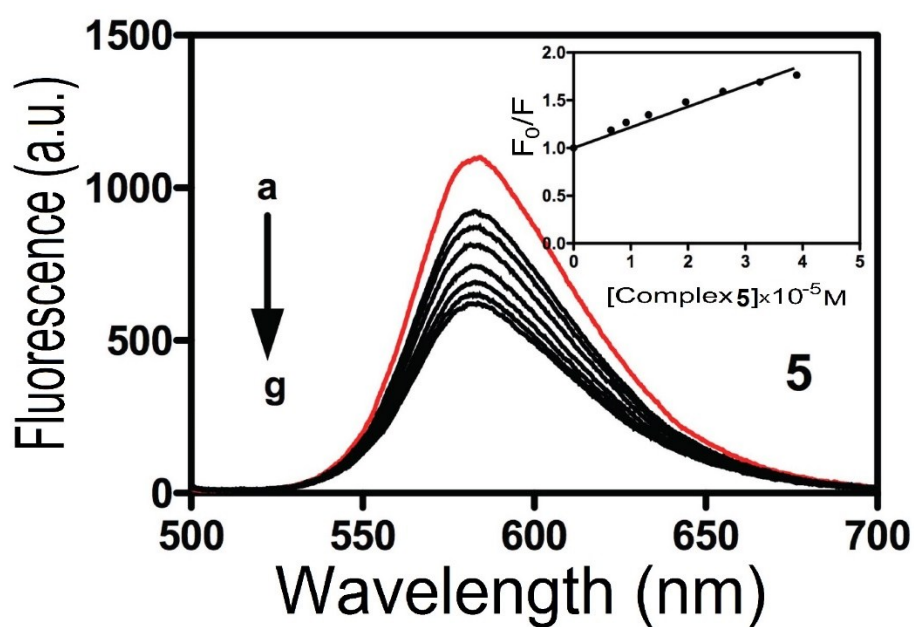
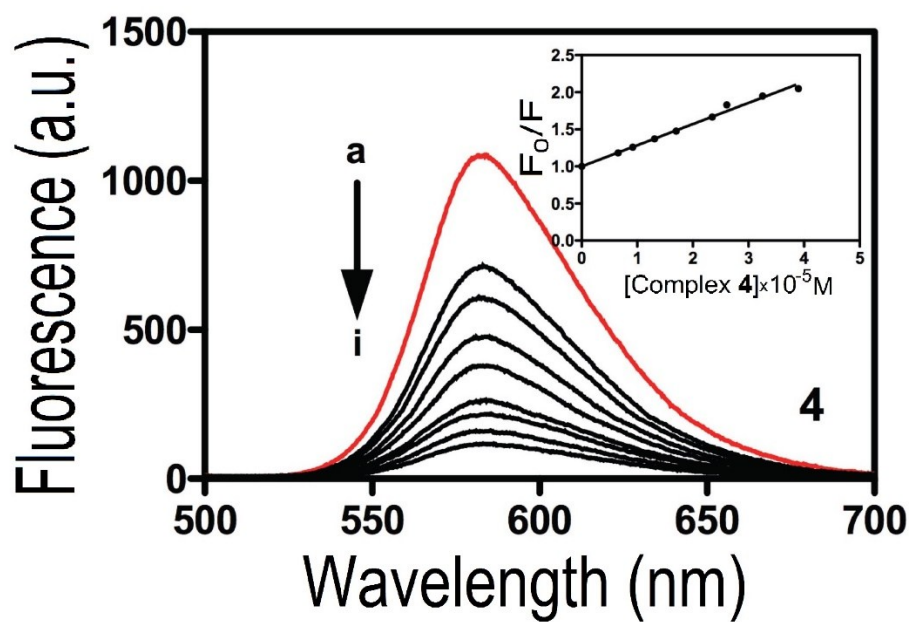
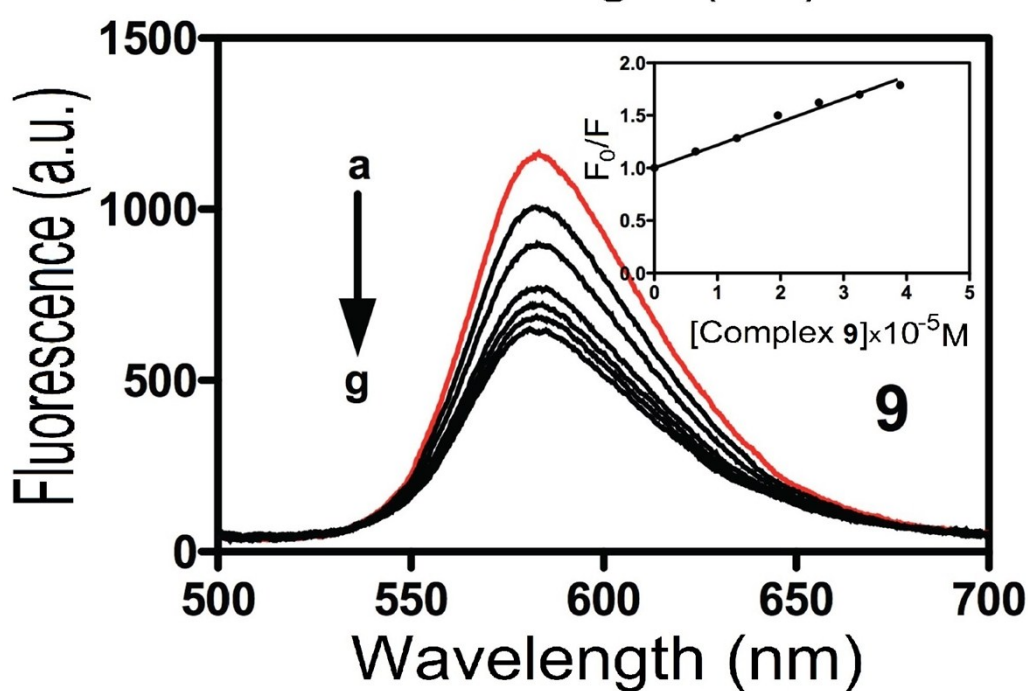
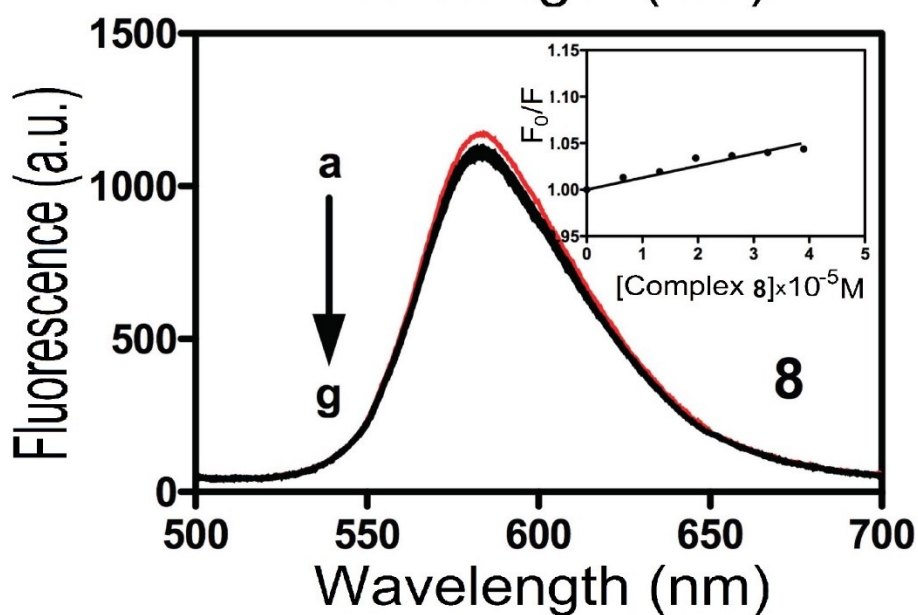
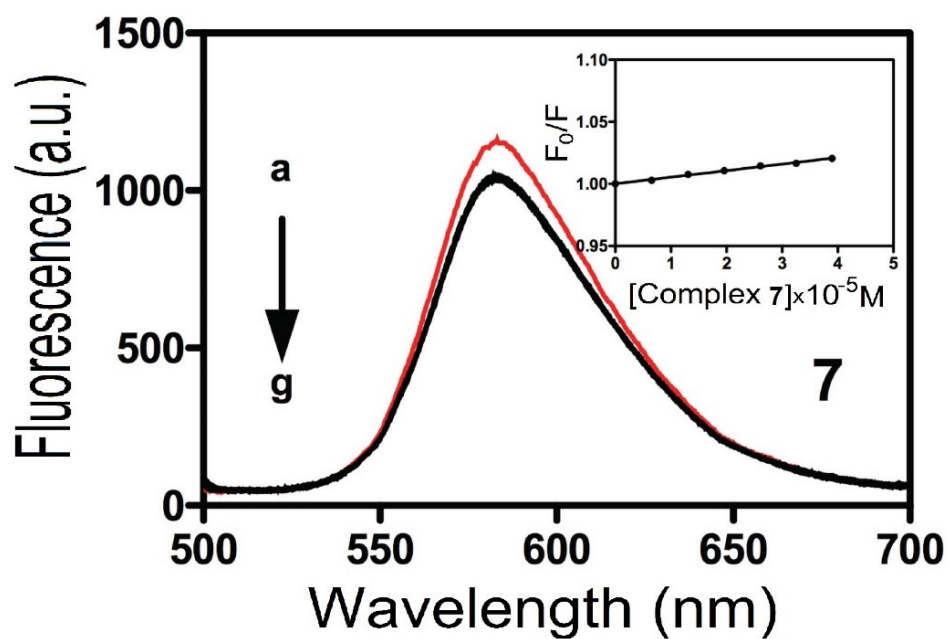


Figure 22S. Absorption spectral traces of different complexes (**panel A**) and ligands (**panel B**) in phosphate buffer (10mM, pH 7.2) after gradual addition of calf thymus DNA (from a = 0 to h = 24 mM). Inset shows the plot of $[DNA]/\epsilon_a - \epsilon_f$ vs. $[DNA]$ (equation 1).







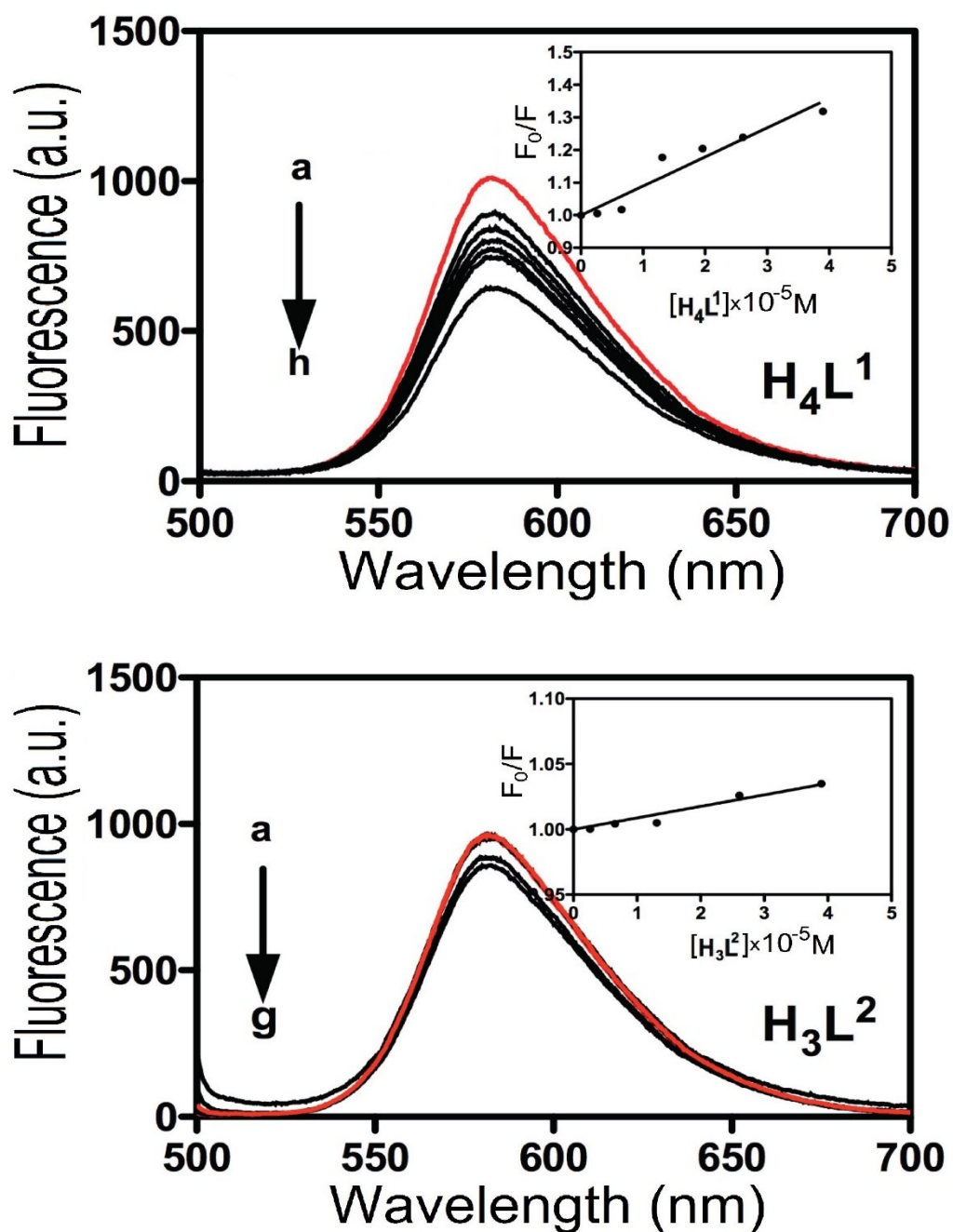
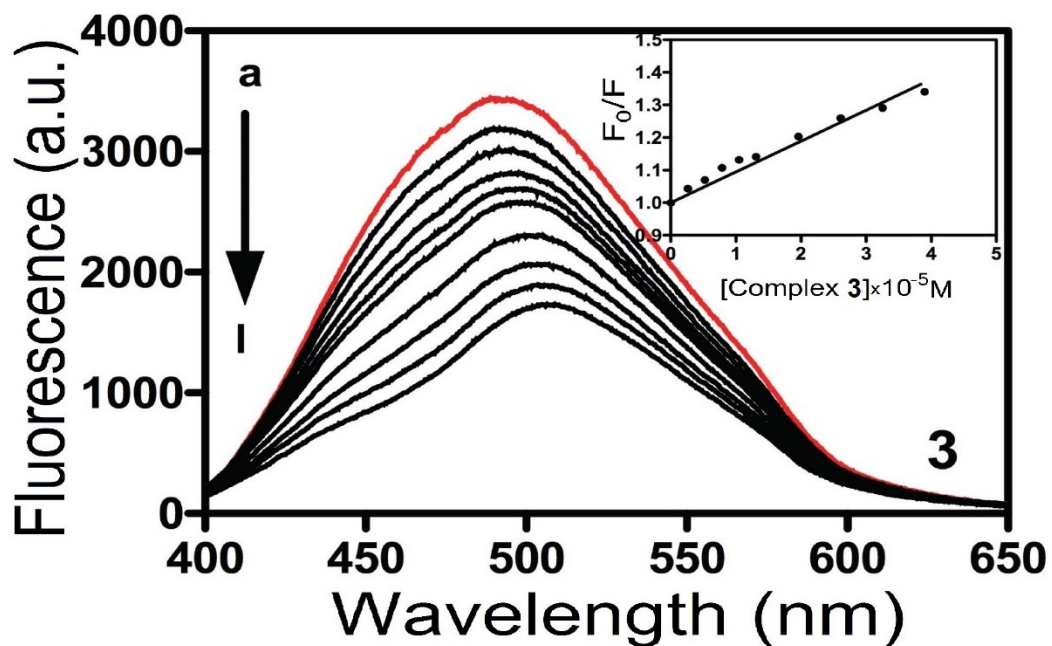
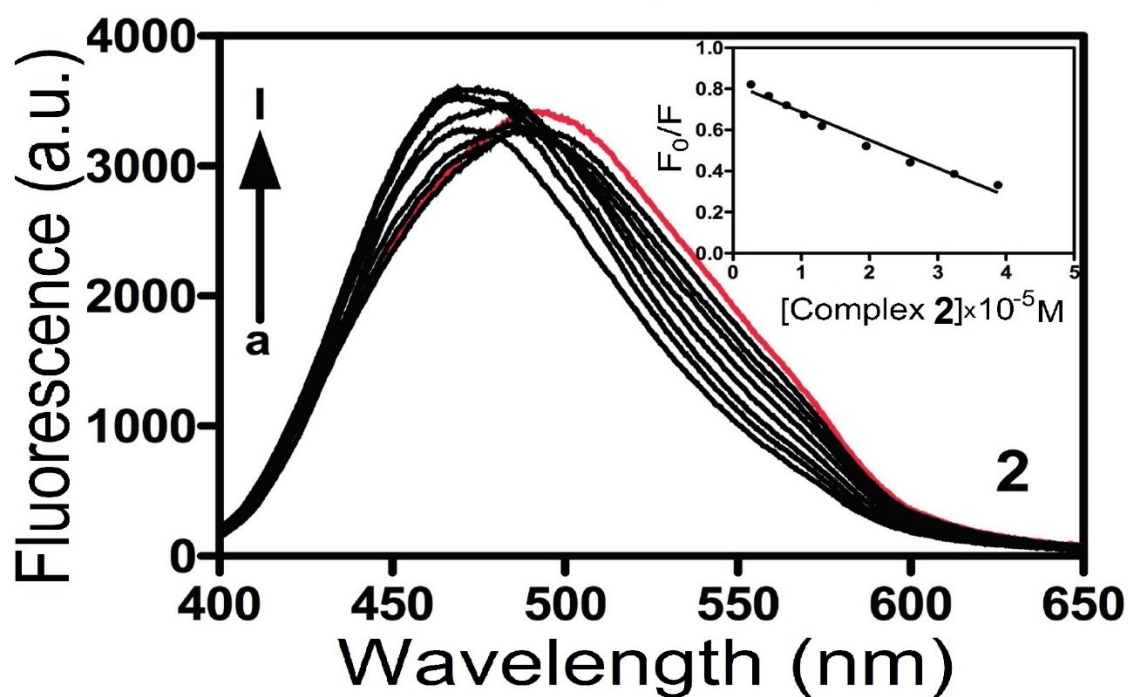
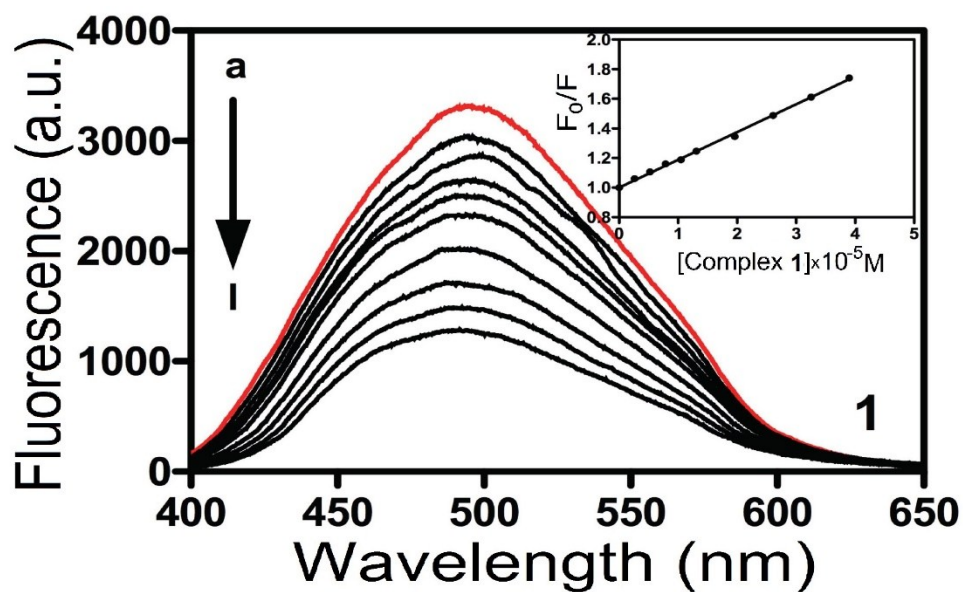
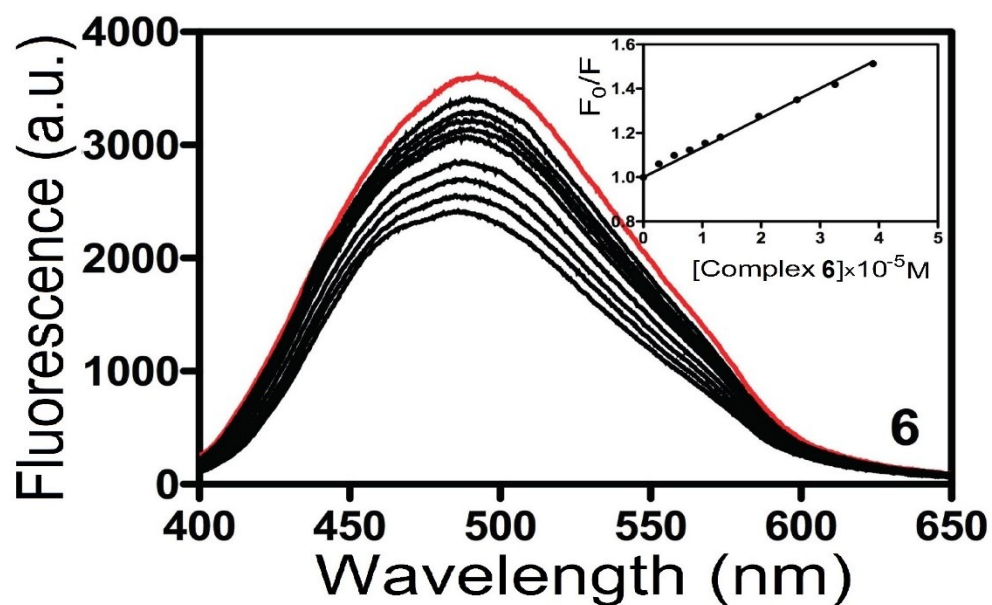
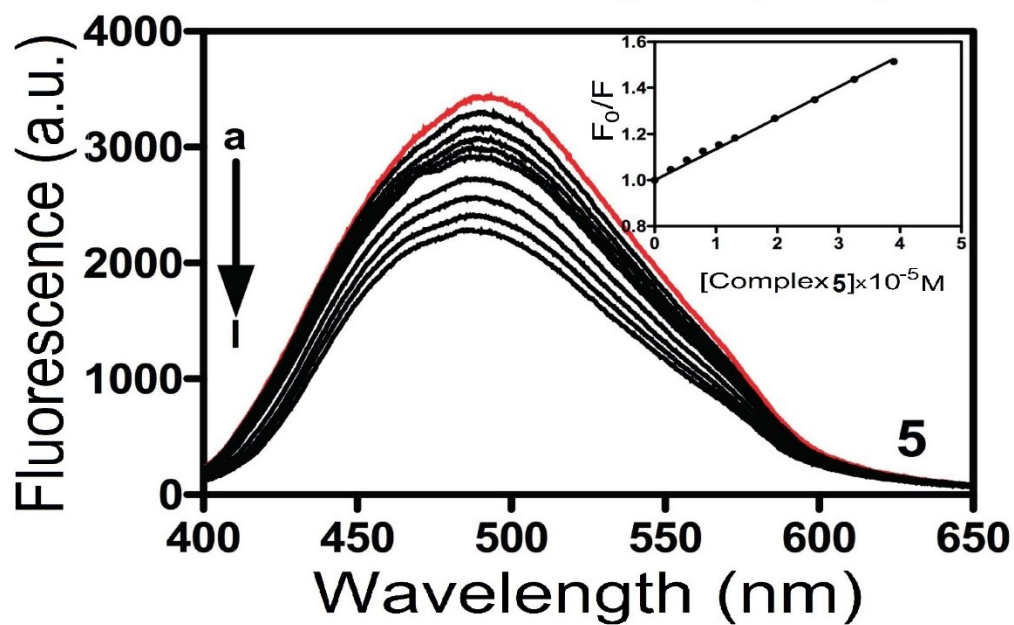
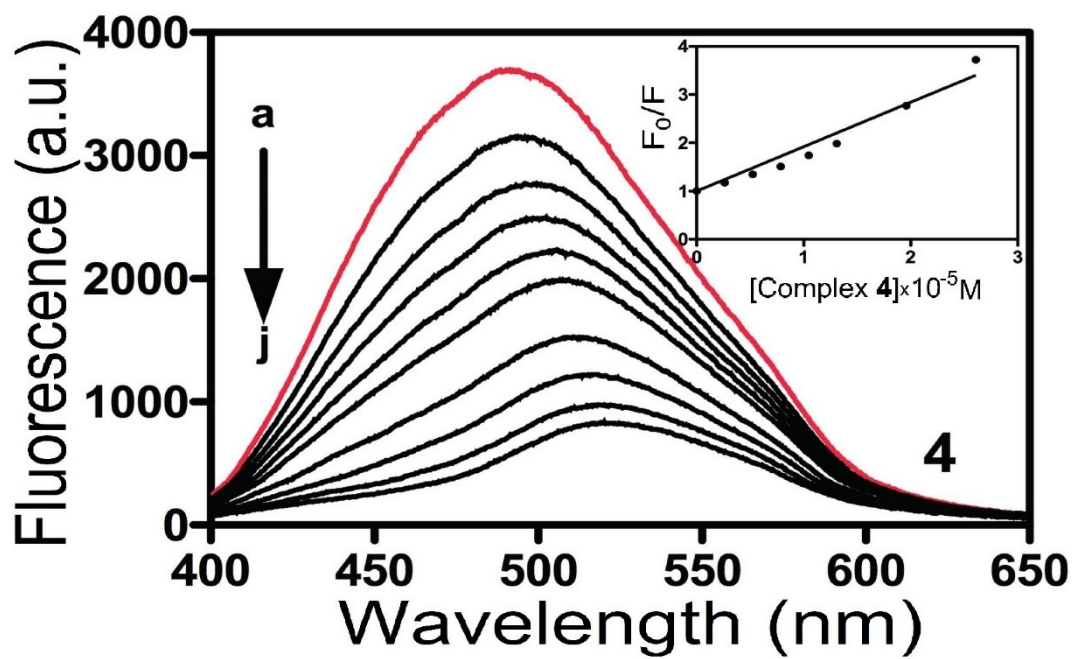
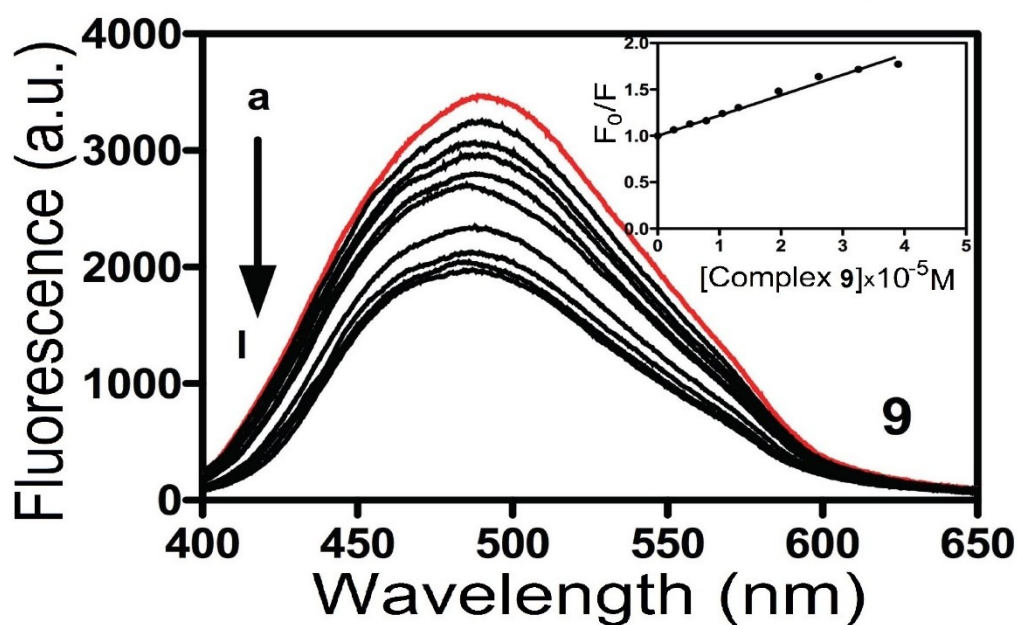
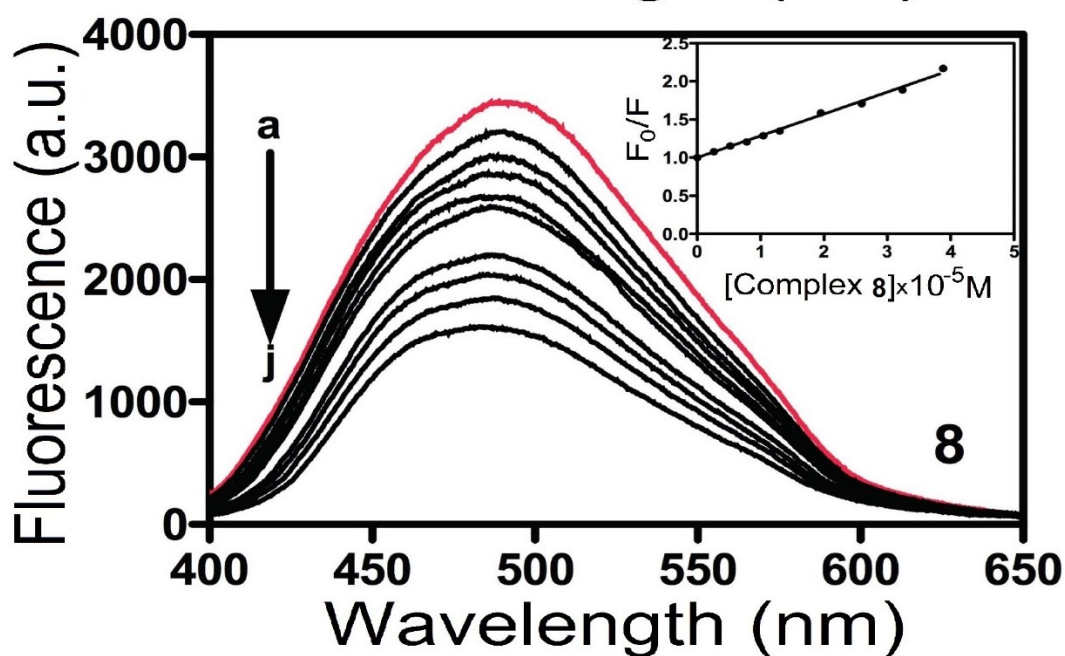
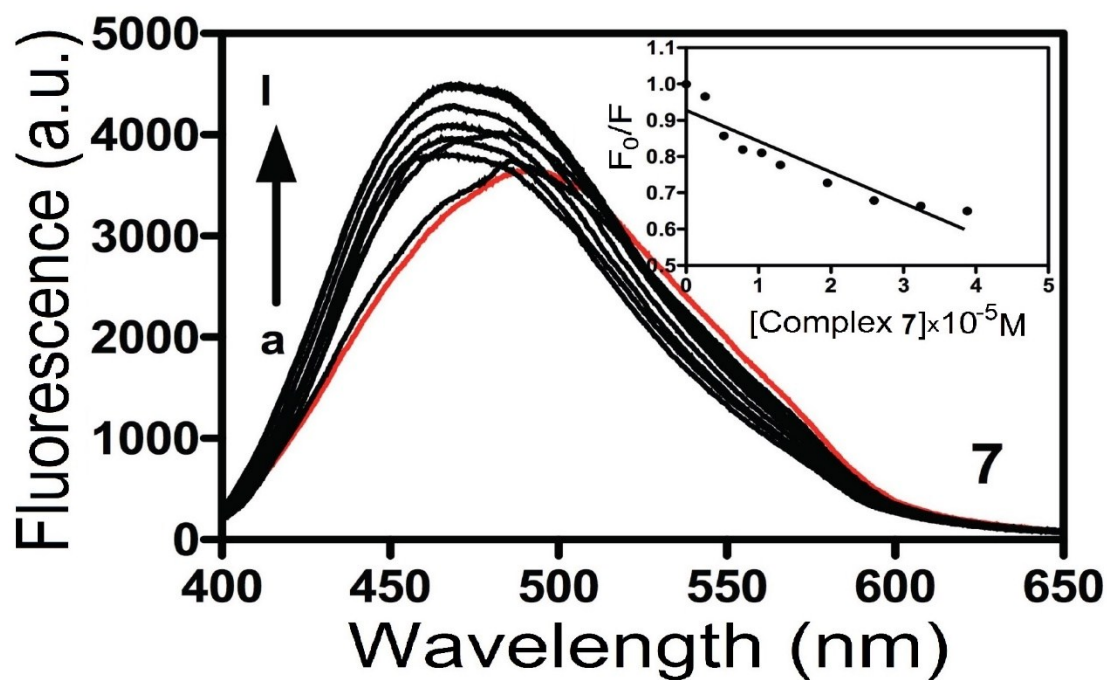


Figure 23S. Emission spectra of EB bound to ct-DNA upon excitation at 490 nm in the presence of increasing concentration of different complexes from (a) 0 to (g) 39 μM ; ($[\text{EB}] = 6 \mu\text{M}$, $[\text{ct-DNA}] = 50 \mu\text{M}$. Inset: Stern Volmer plot of F_0/F vs. $[\text{complex}]$ for the titration of the complexes to EB-ct-DNA complex.







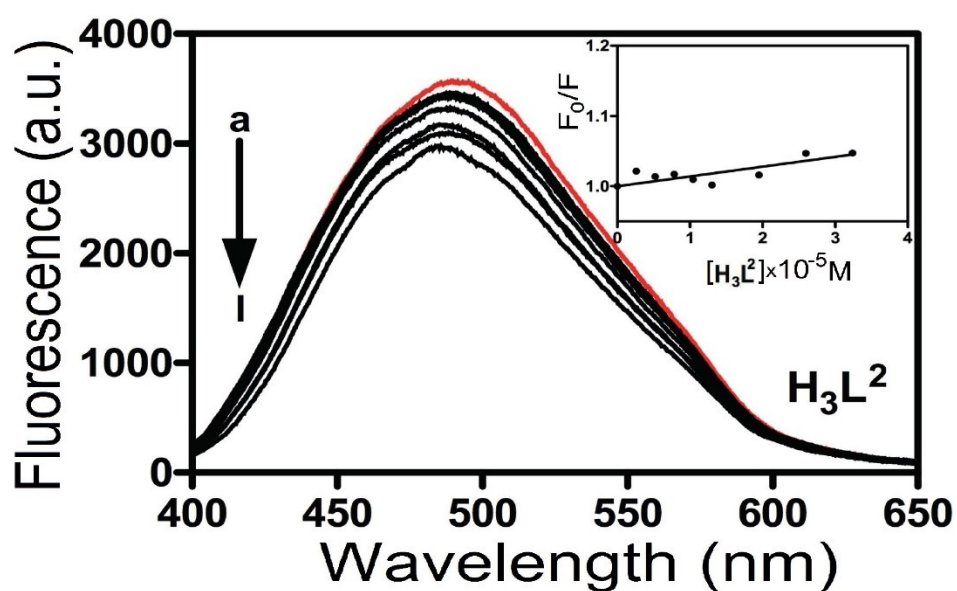
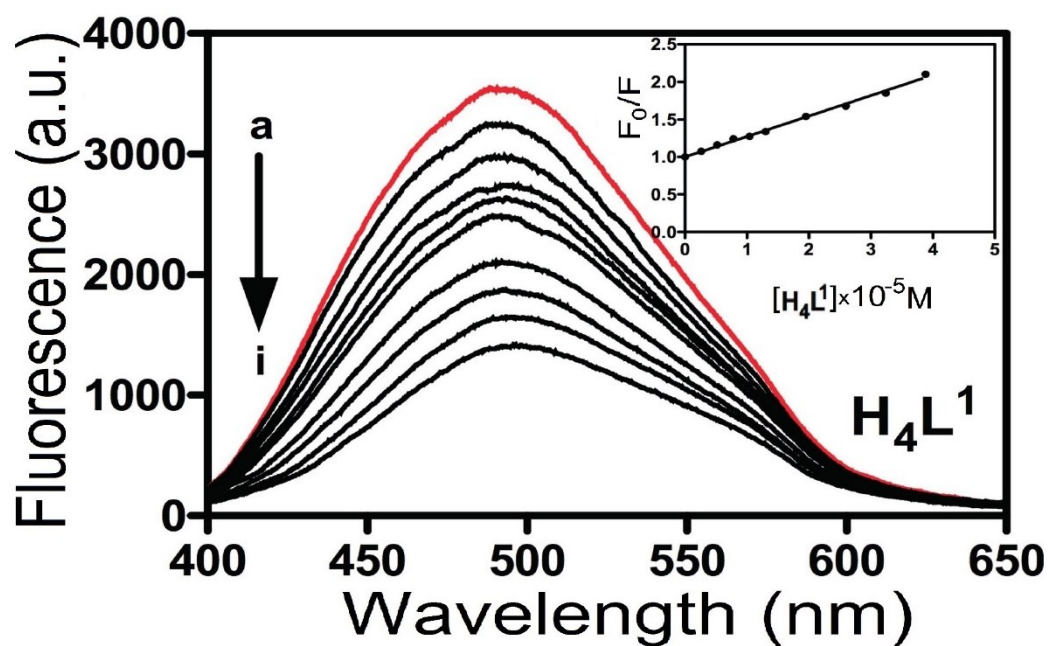
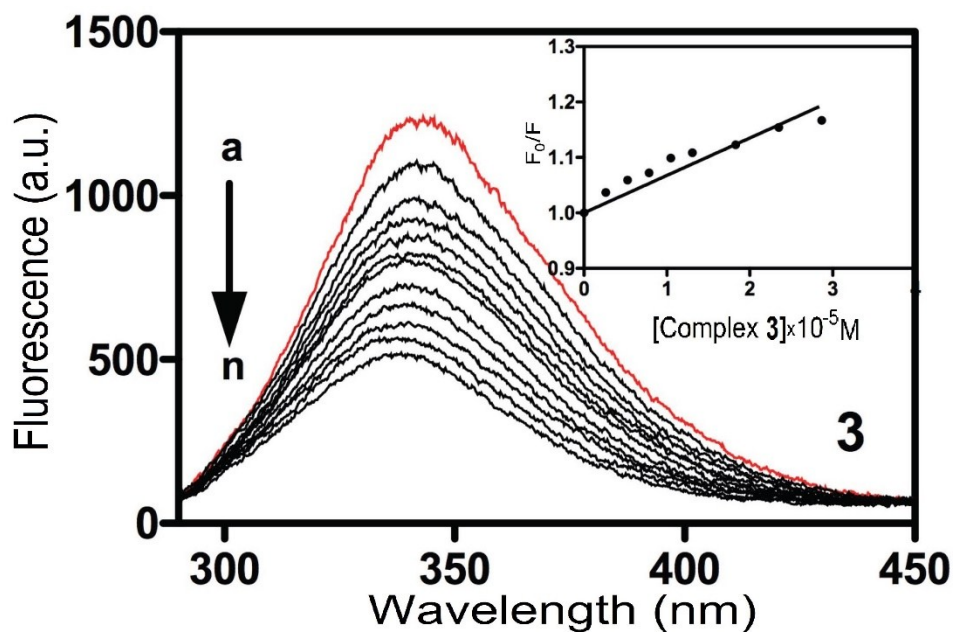
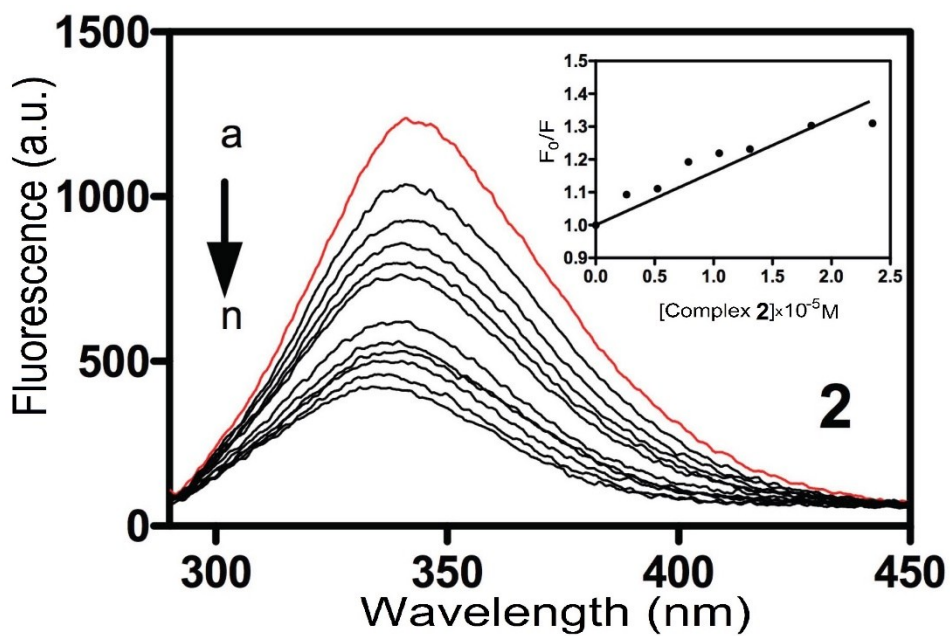
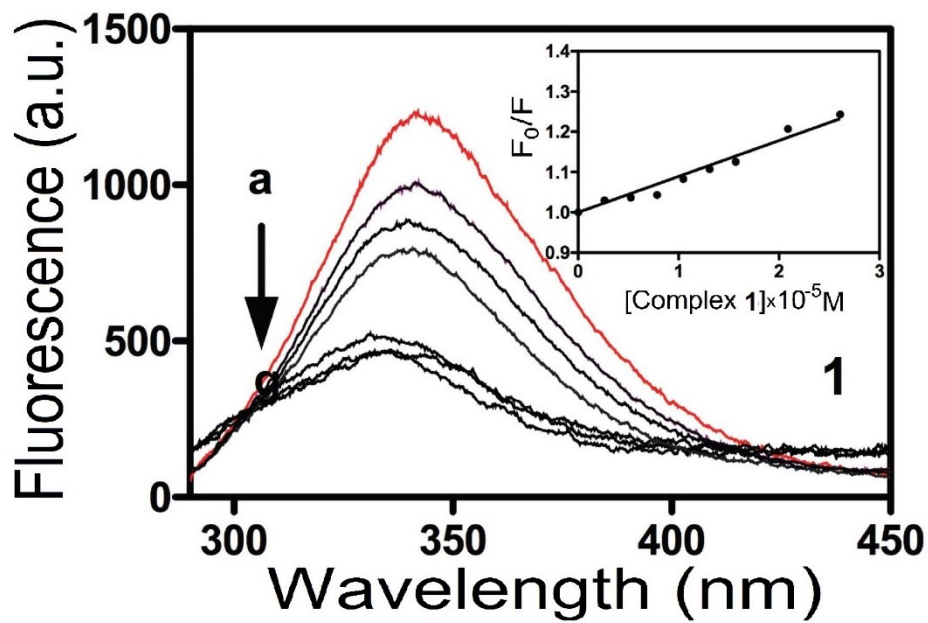
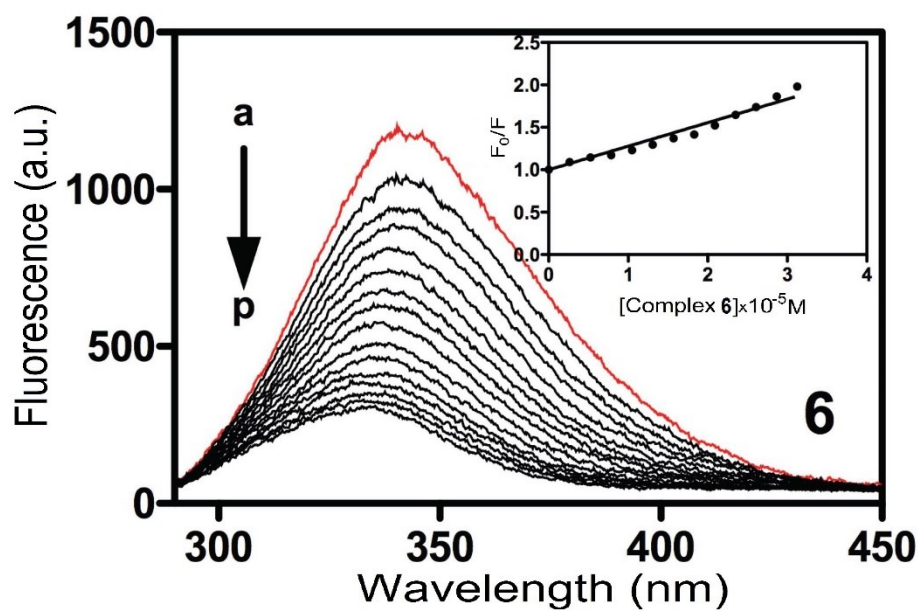
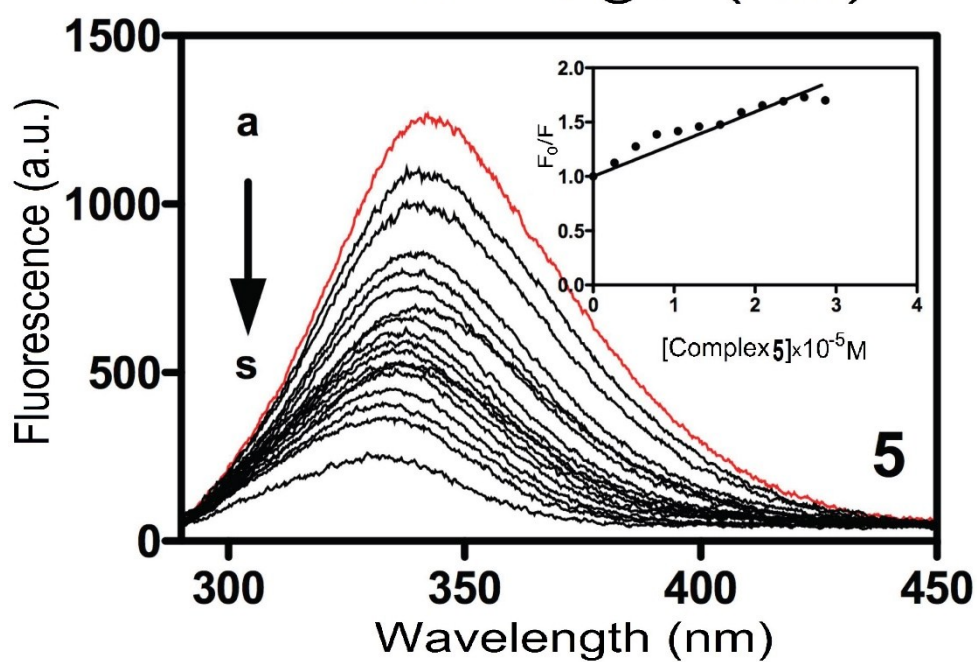
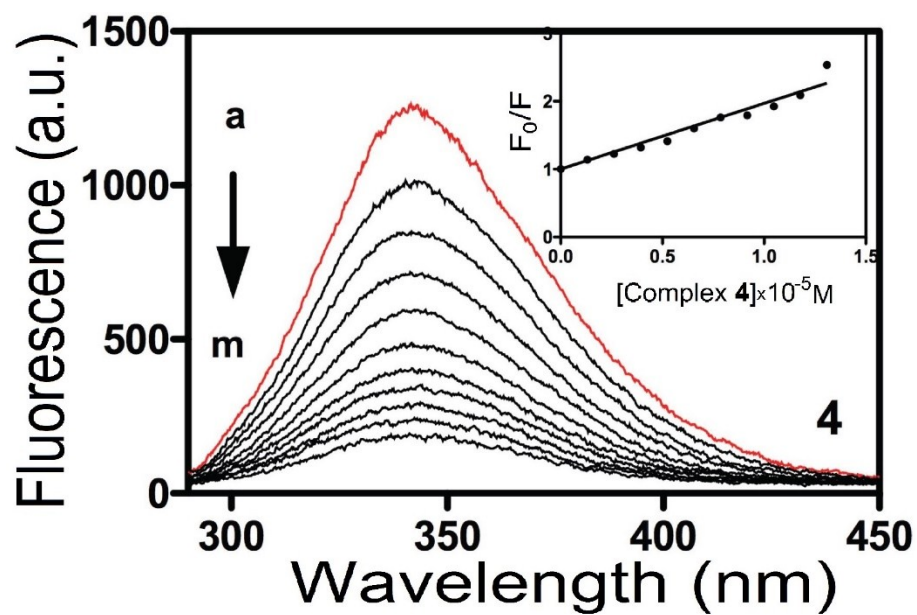
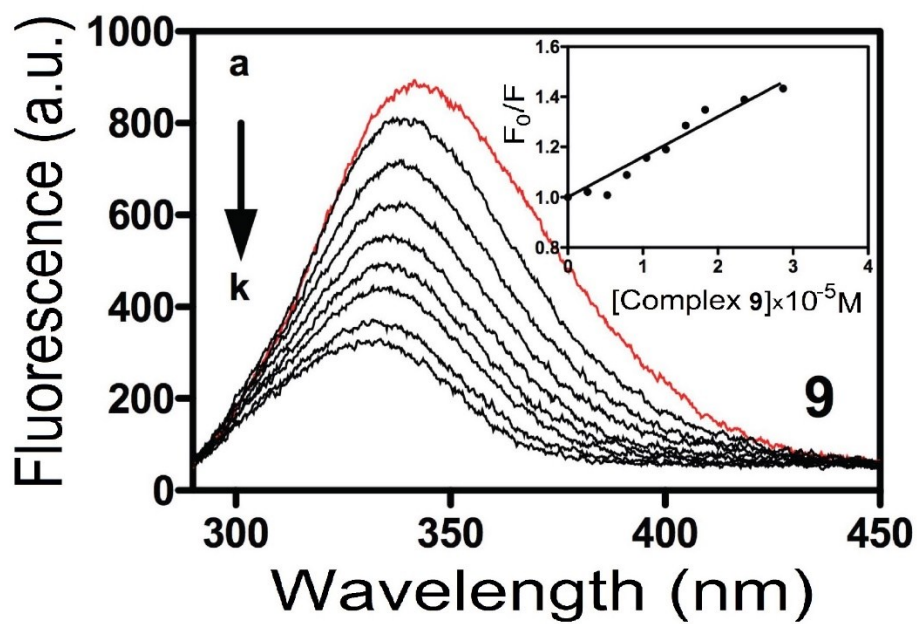
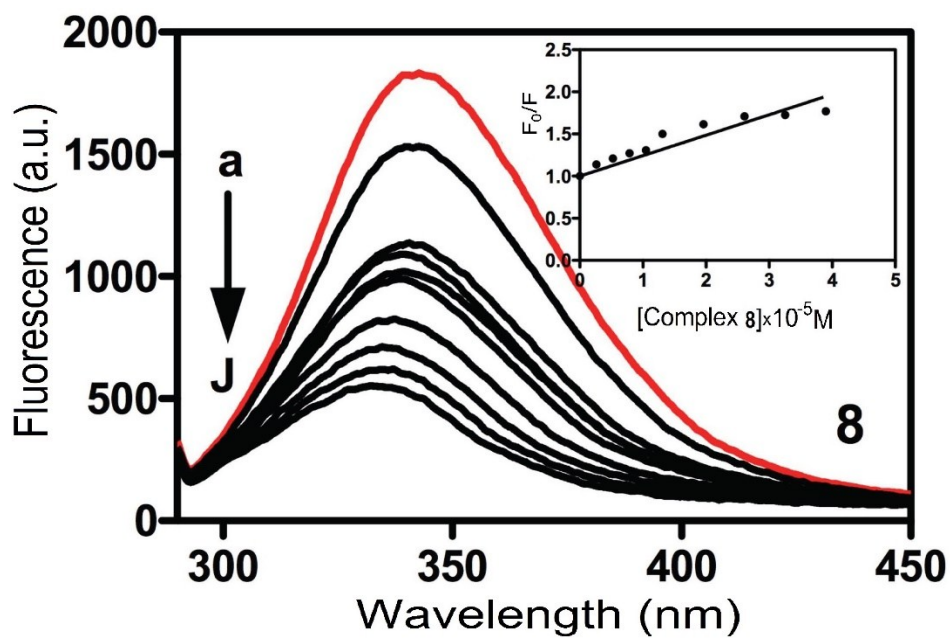
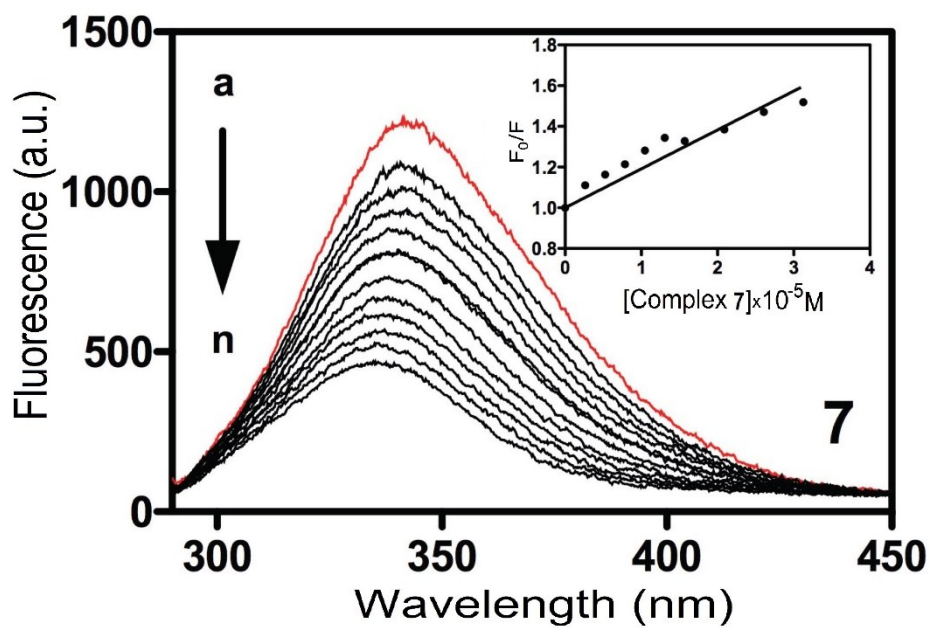


Figure 24S. Emission fluorescence spectra of DAPI-ct-DNA complex upon excitation at 338 nm in the presence of increasing concentration of different compounds and proligands in the range 0-36 μM . Experiments were performed in 10 mM phosphate buffer, pH 7.2. Inset: Stern-Volmer plot of F_0/F vs. [complex] for the titration of the complexes to DAPI-ct-DNA complex







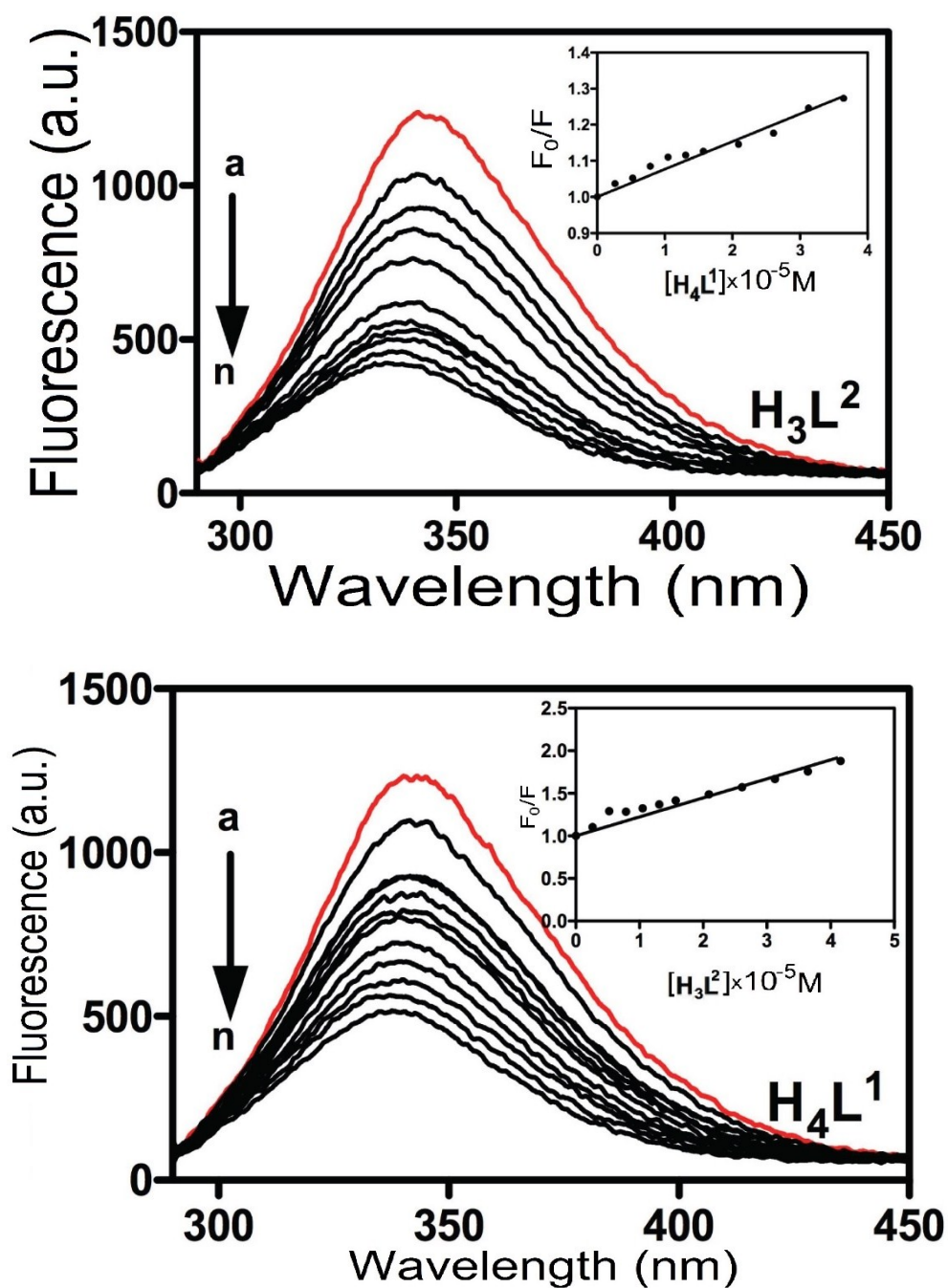


Figure 25S. Fluorescence emission spectra of BSA upon excitation at 280 nm, in the absence and presence of different concentrations of compounds and proligands. Inset, Stern-Volmer plots of F_0/F vs. [complex 4] for the titration of the complexes to BSA.