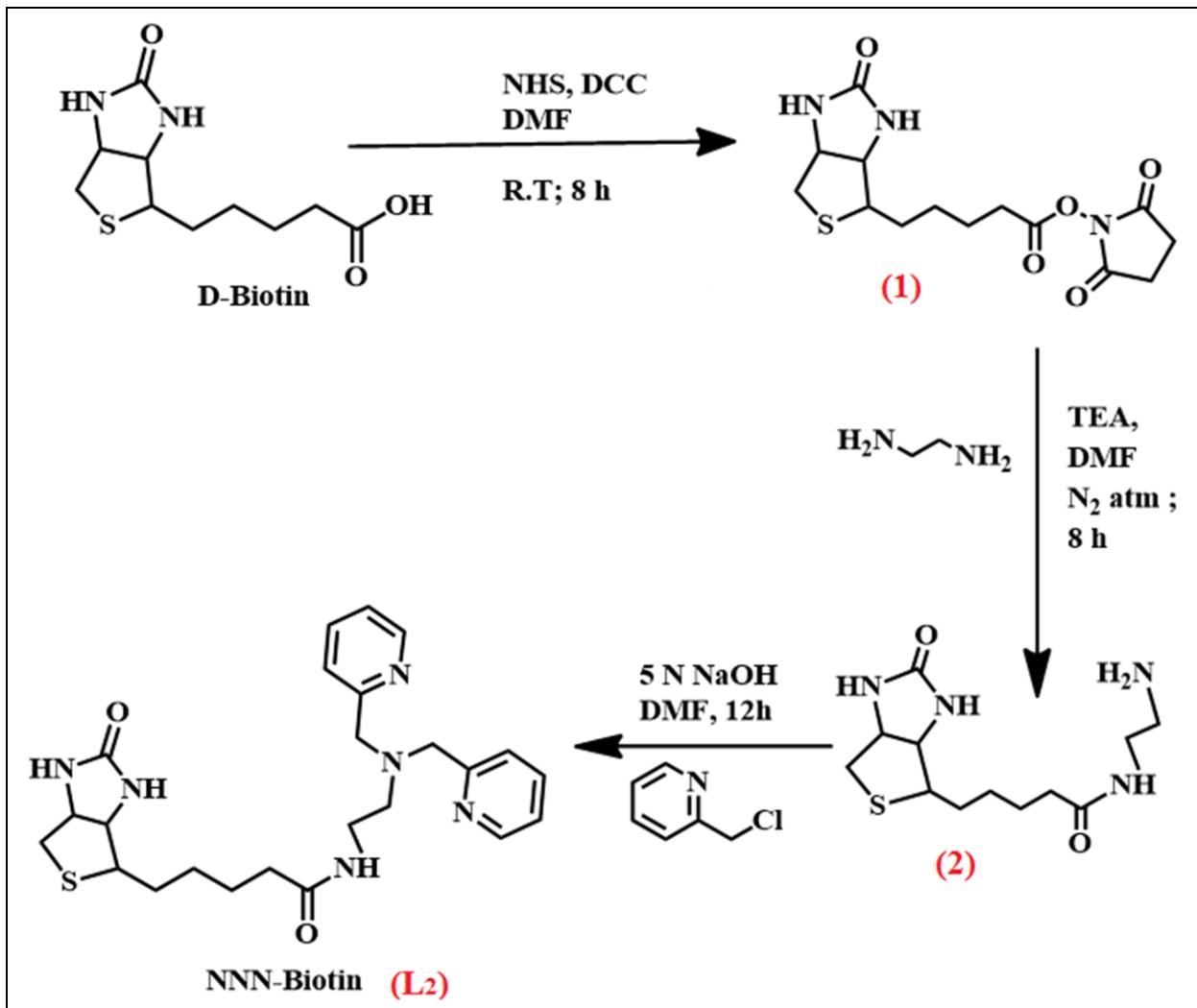


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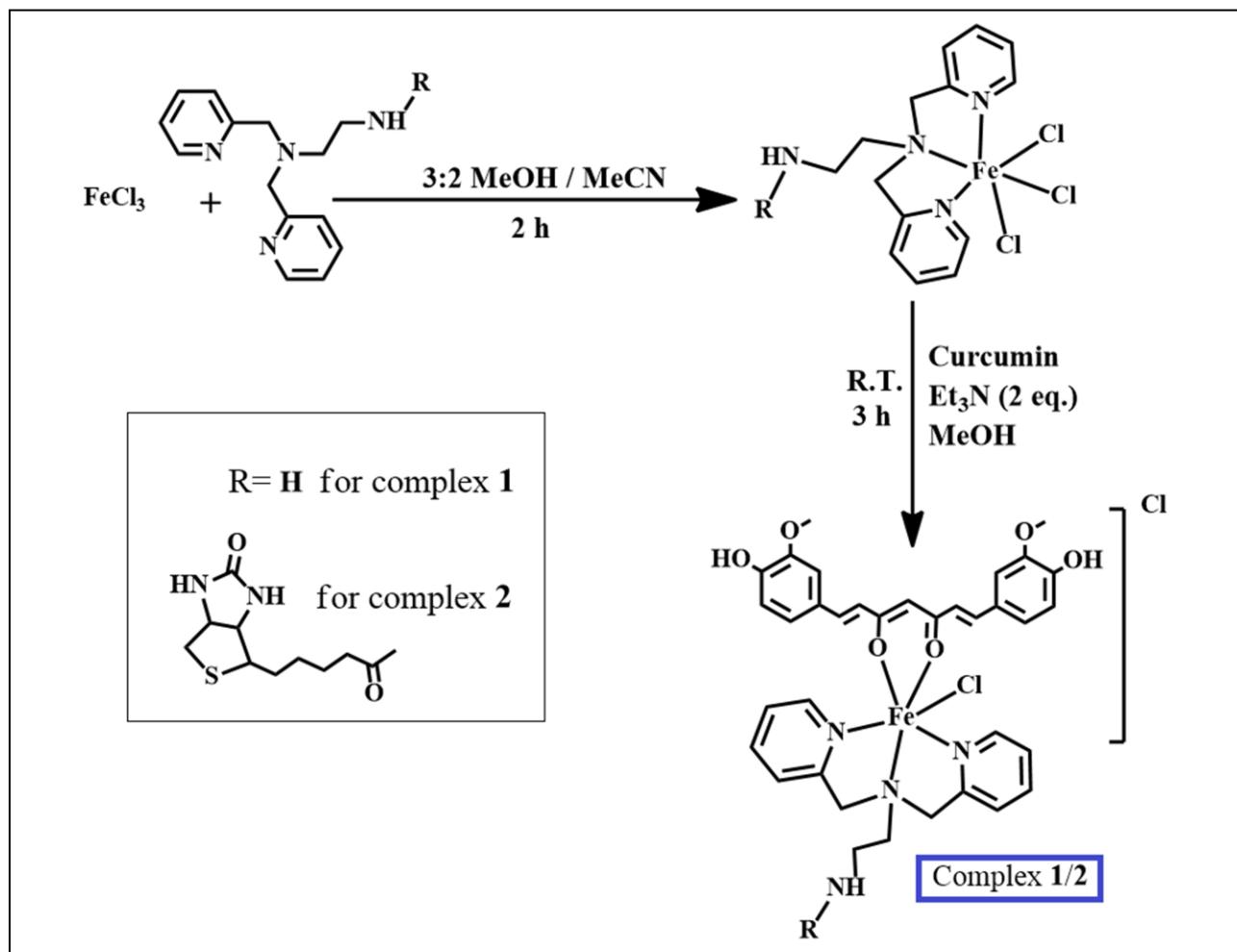
## **Supporting Information**

### **Biotin-Appended Iron(III) Complexes of Curcumin for Targeted Photo-Chemotherapy**

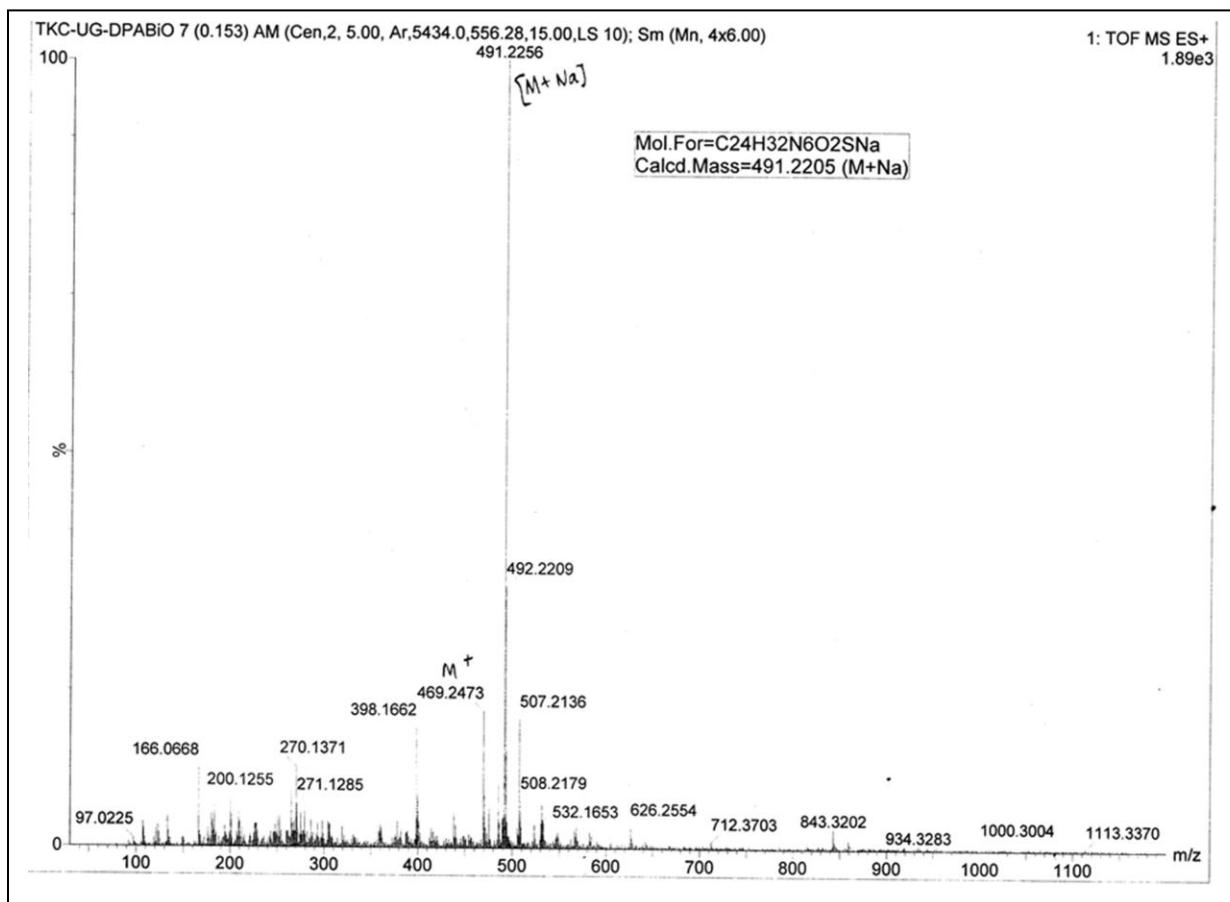
Somarupa Sahoo, Abinaya Raghavan, Arun Kumar, Dipankar Nandi,\* and Akhil R. Chakravarty\*



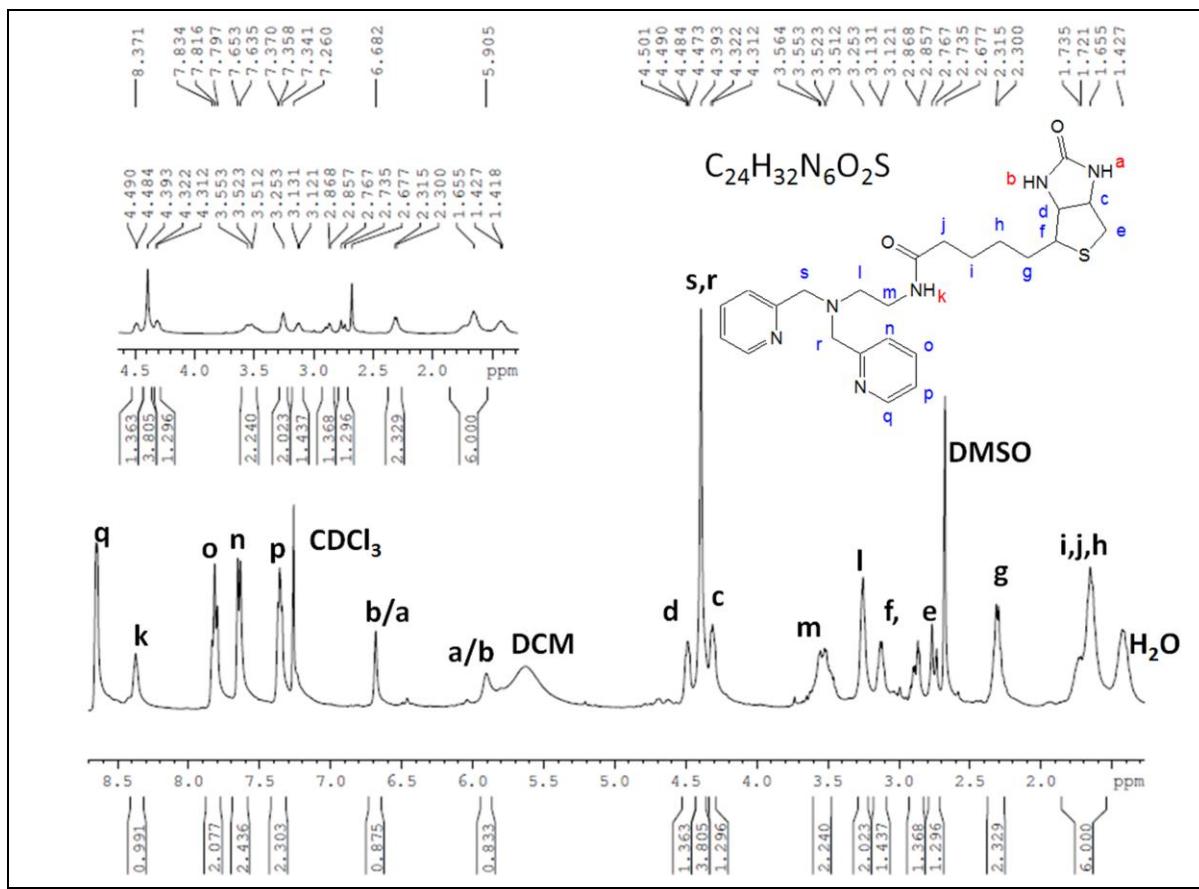
**Scheme S1.** Synthetic scheme for the preparation of *NNN*-donor biotin-appended dipicolylamine ligand ( $L^2$ ).



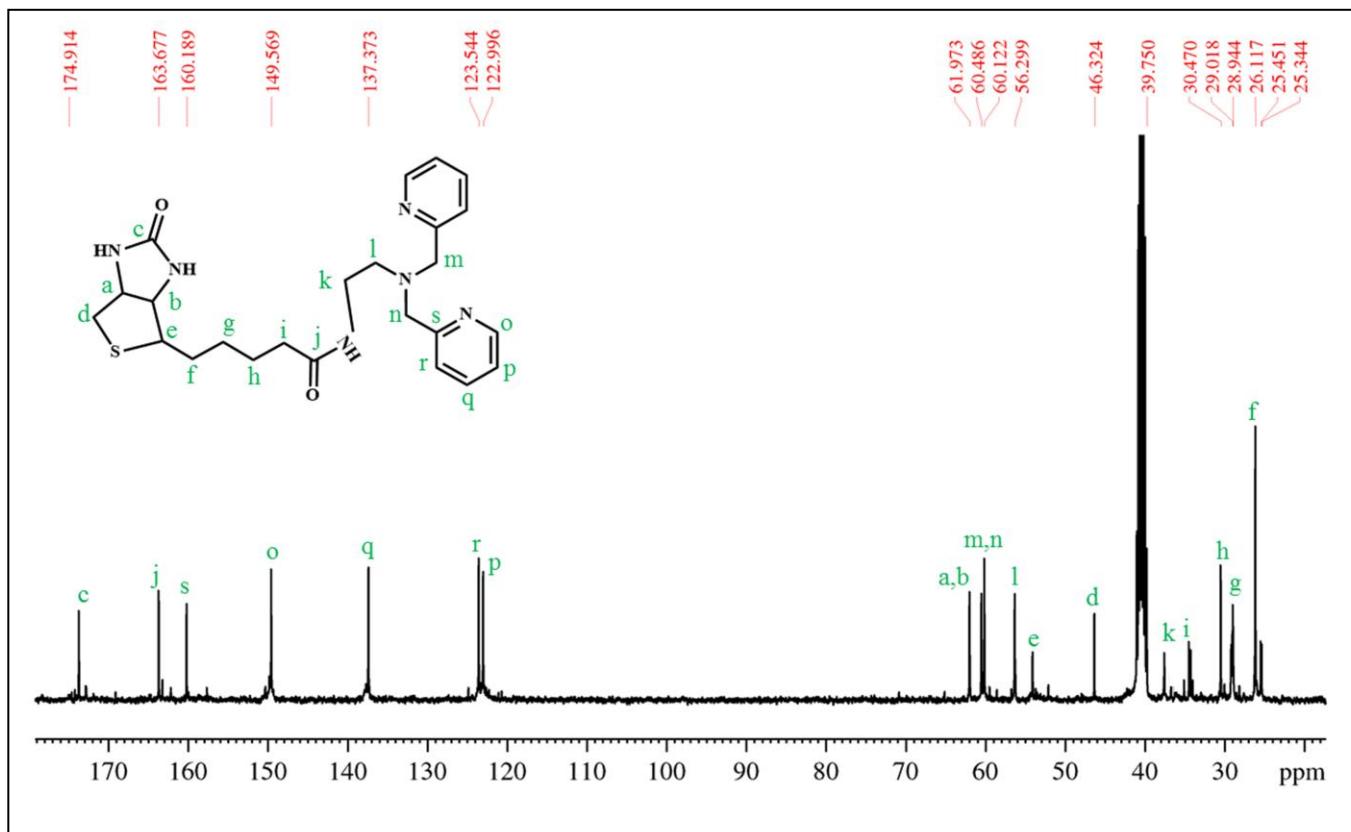
**Scheme S2.** Generalized synthetic scheme for the iron(III) complexes **1** and **2**.



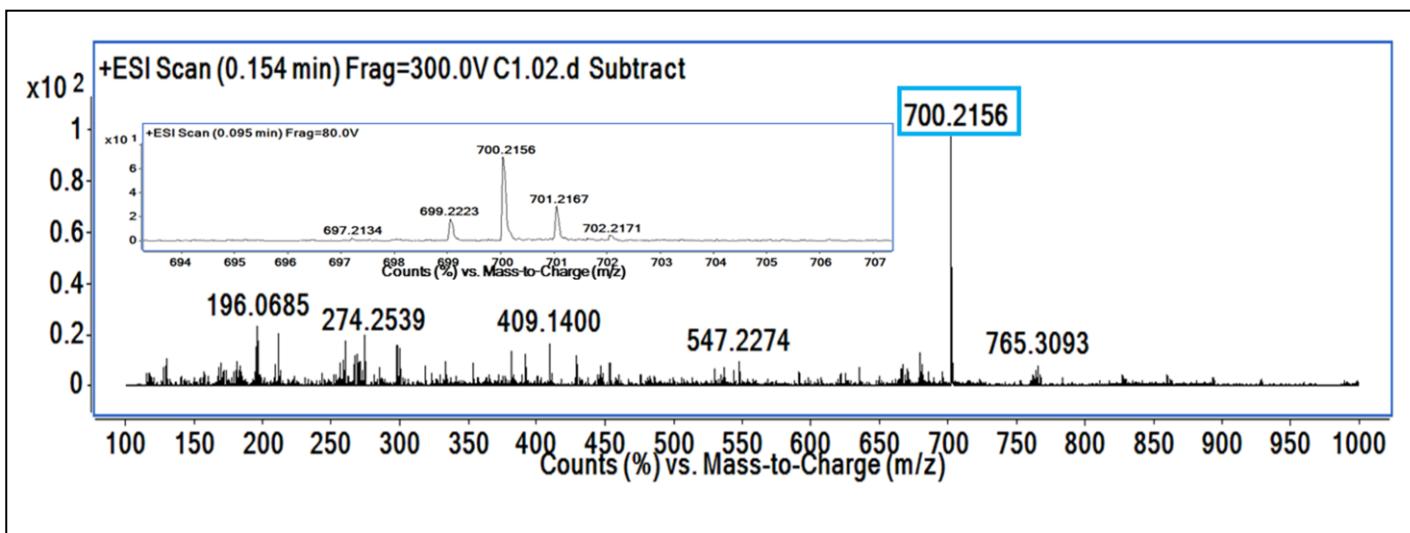
**Figure S1.** ESI-MS spectrum of the biotin ligand ( $L^2$ ) in MeOH showing a prominent peak corresponding to  $[M + Na]^+$  at 491.2256 (m/z).



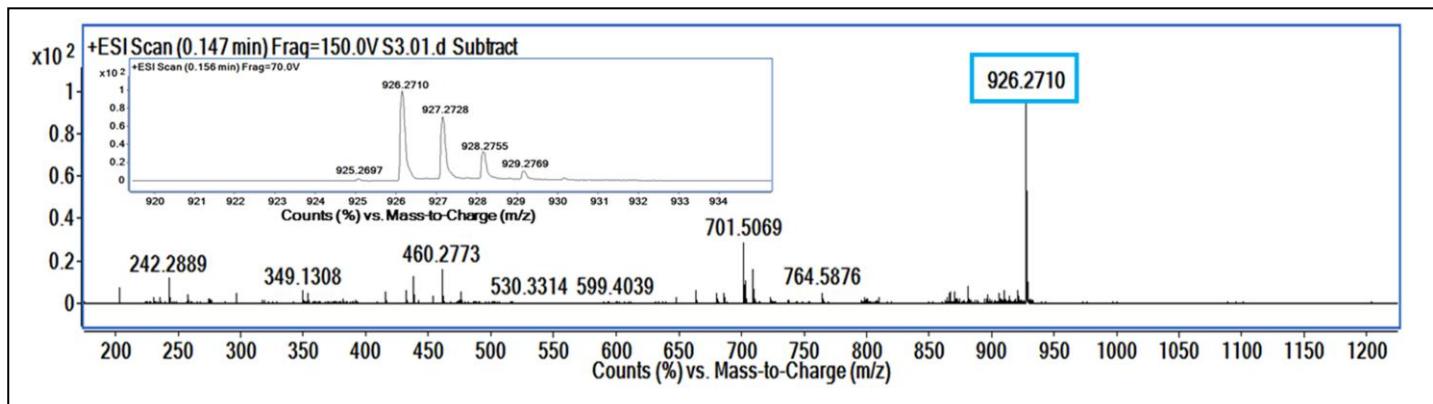
**Figure S2.**  $^1\text{H}$  NMR spectrum of ligand L<sup>2</sup> in  $\text{CDCl}_3$ .



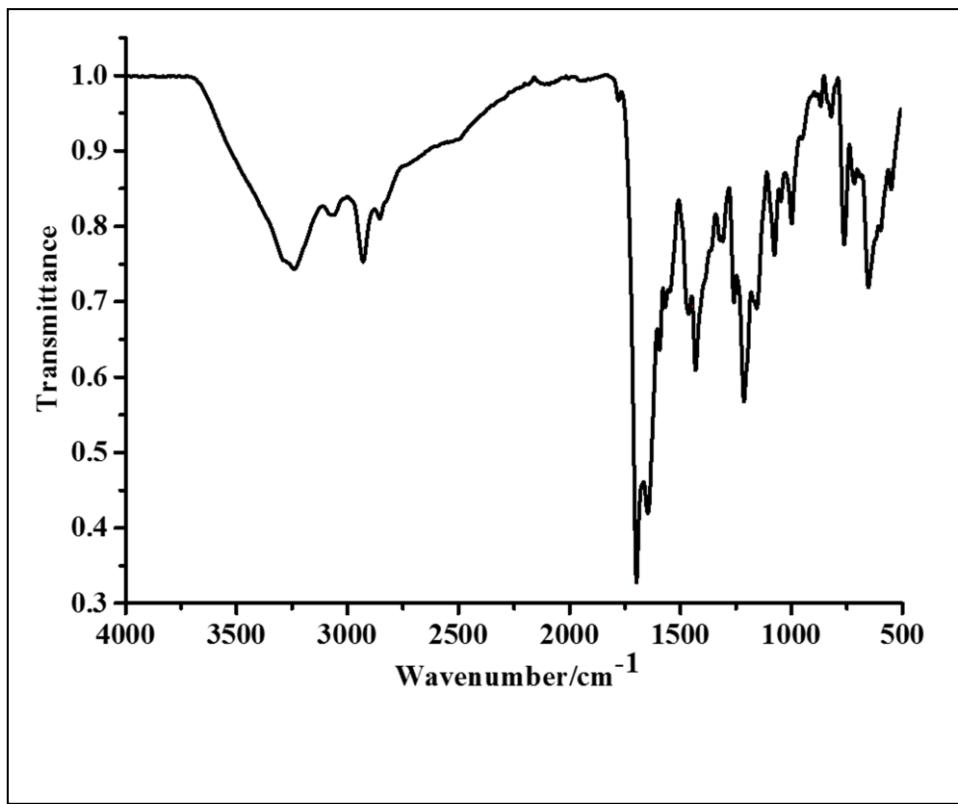
**Figure S3.**  $^{13}\text{C}$  NMR spectrum of ligand  $\text{L}^2$  in  $\text{DMSO-d}_6$ .



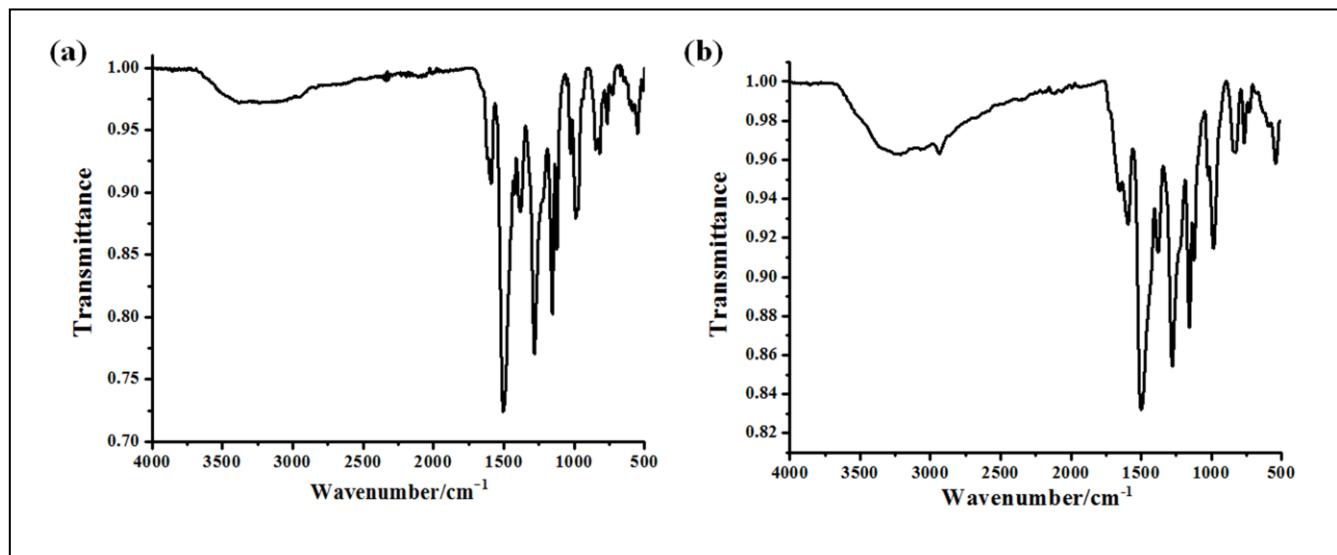
**Figure S4.** ESI-MS spectrum of complex **1** in acetonitrile showing a prominent peak corresponding to  $[\text{M}-\text{Cl}]^+$  at  $700.2156$  ( $m/z$ ). Inset shows the isotopic distribution pattern.



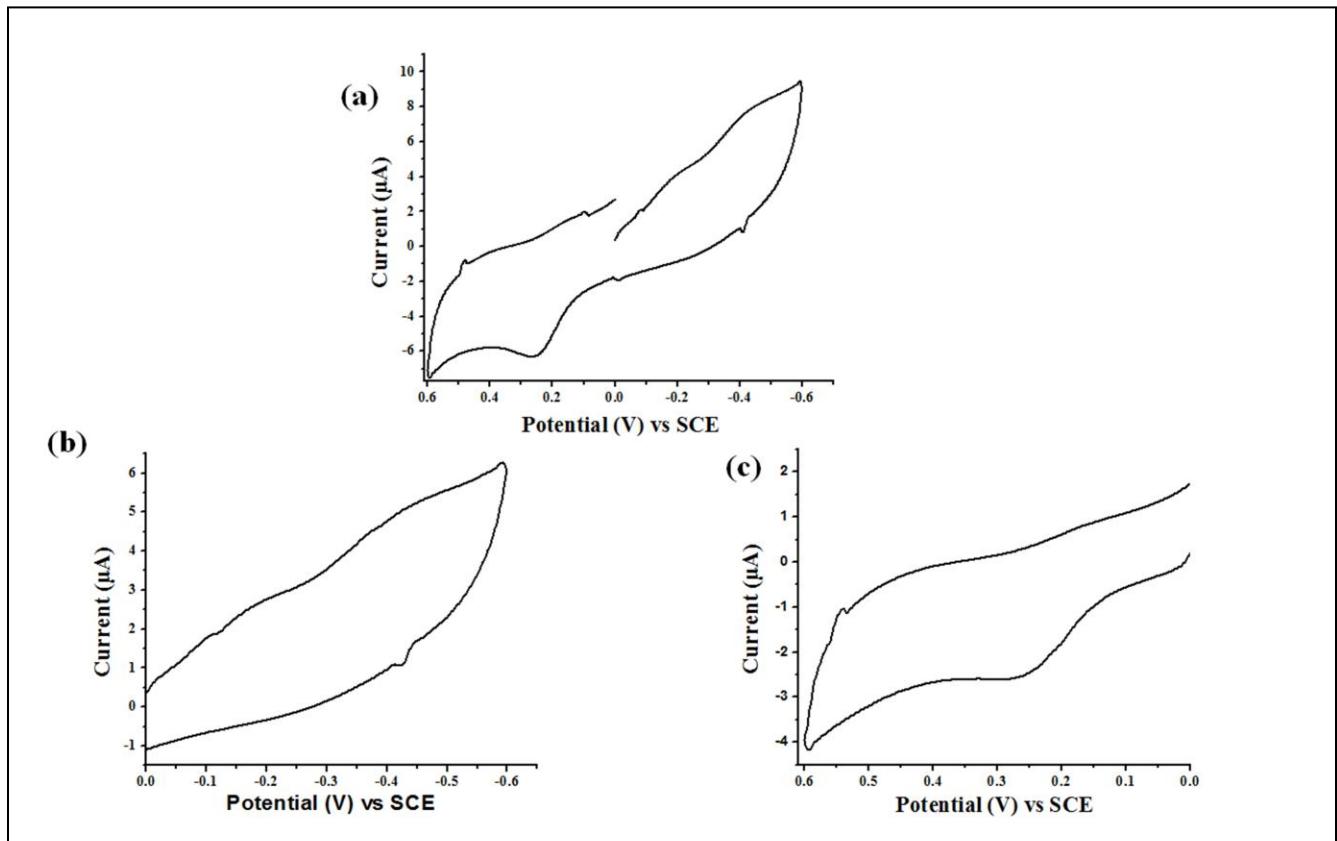
**Figure S5.** ESI-MS spectrum of complex **2** in acetonitrile showing a prominent peak corresponding to  $[M-Cl]^+$  at 926.2710 ( $m/z$ ). Inset shows the isotopic distribution pattern.



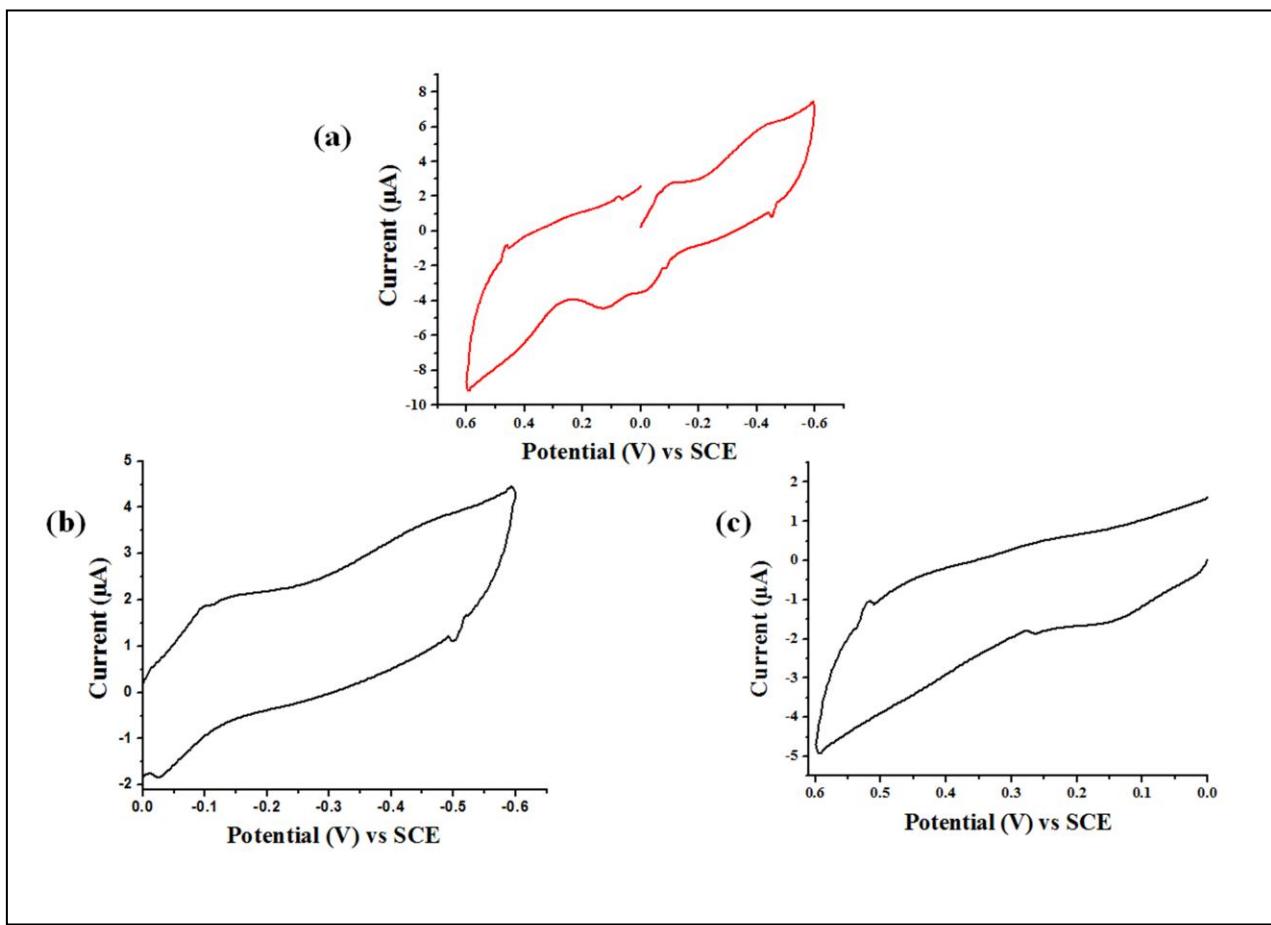
**Figure S6.** IR spectrum of ligand  $L^2$  in the solid phase showing a strong peak for keto the group observed at  $1660\text{ cm}^{-1}$ .



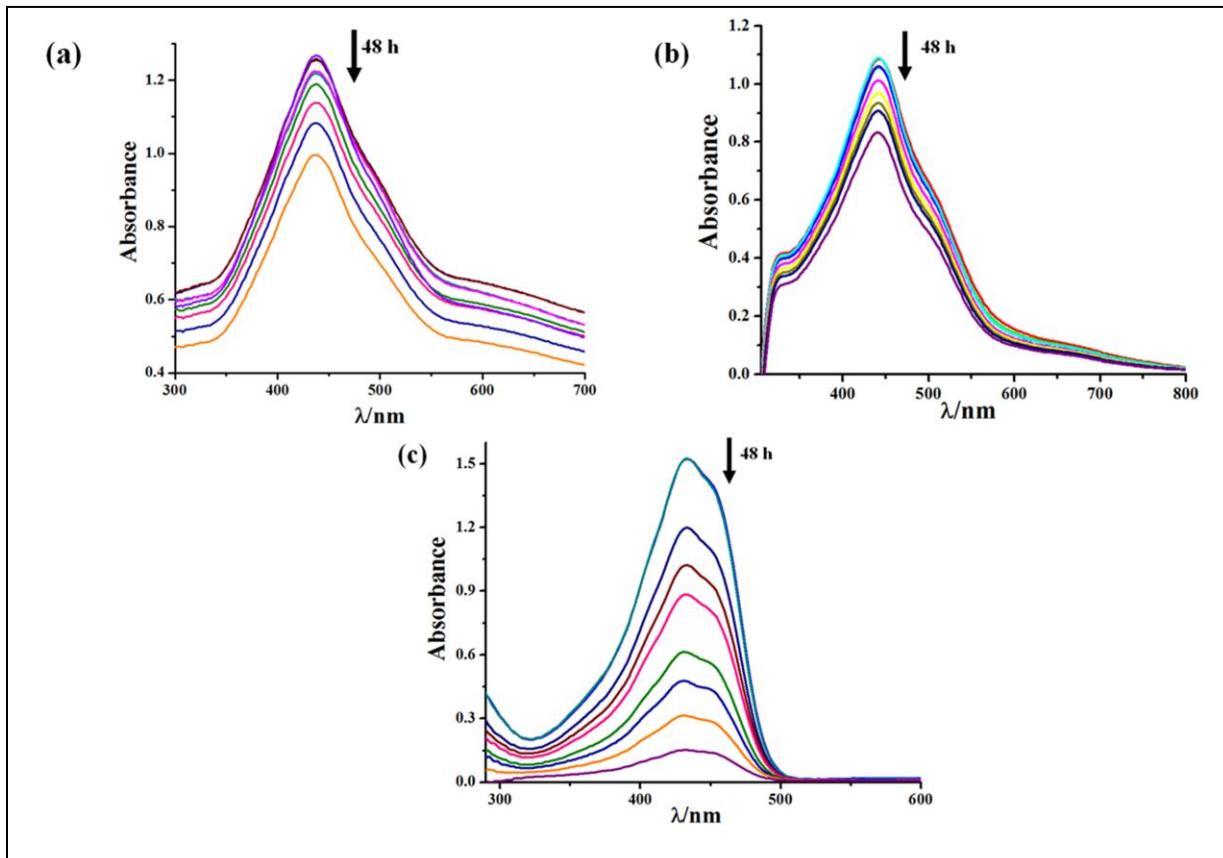
**Figure S7.** IR spectra of the complexes **1** and **2** in the solid phase: **1**, (a); **2**, (b).



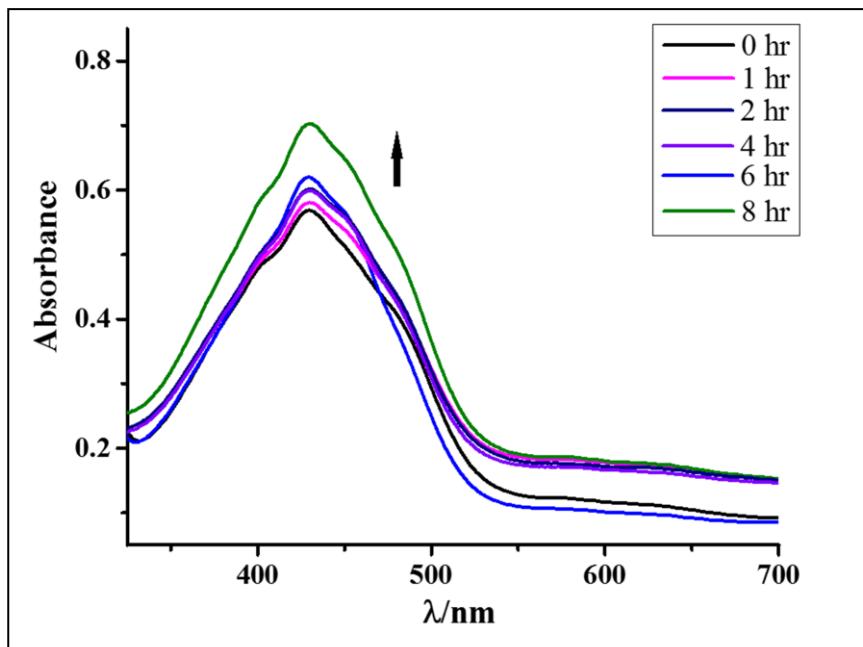
**Figure S8.** Cyclic voltammograms of complex **1** showing the Fe(III)/Fe(II) redox response in 0.1 M TBAP-DMF at a scan rate of 100 mV s<sup>-1</sup> : Full scan, (a); Cathodic scan, (b); Anodic scan, (c). The redox responses are sluggish in nature without showing any reversibility.



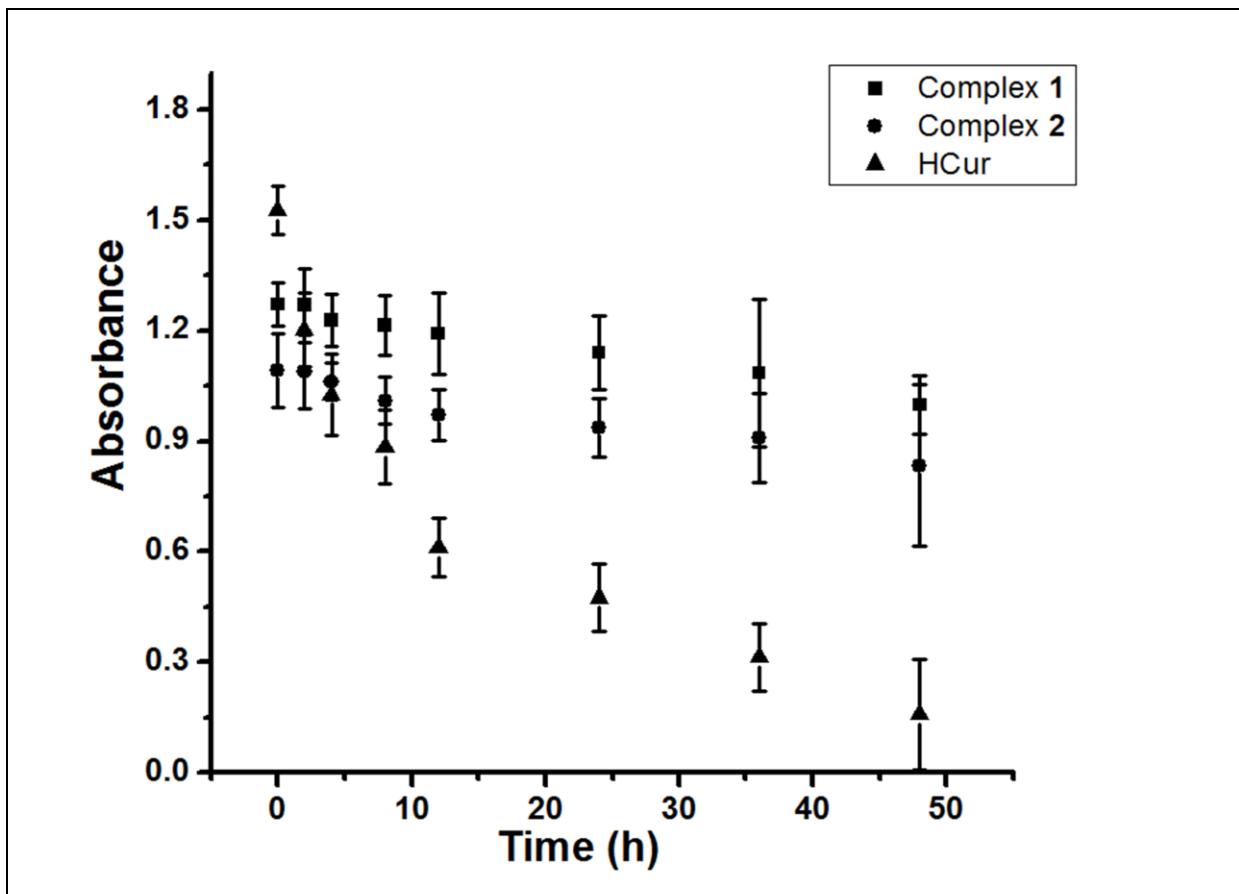
**Figure S9.** Cyclic voltammograms of complex **2** showing the Fe(III)/Fe(II) redox responses in 0.1 M TBAP-DMF at a scan rate of  $100 \text{ mV s}^{-1}$  : Full scan, (a); Cathodic scan, (b); Anodic scan, (c). The redox responses are sluggish in nature.



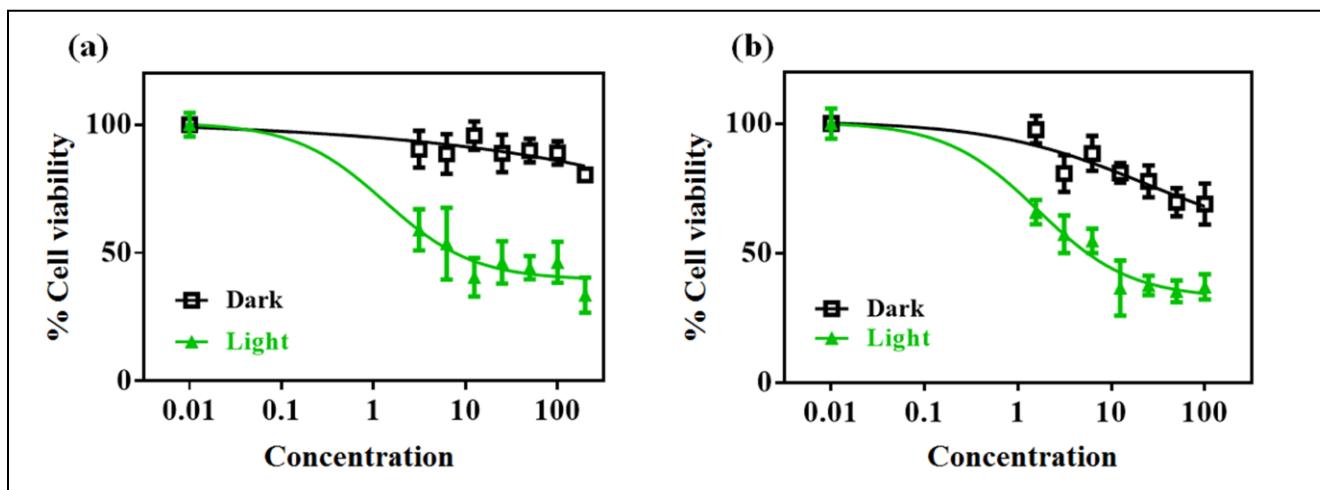
**Figure S10.** The UV-visible spectral traces of the complexes in DMSO/DPBS (1:1 v/v) recorded over a period of 48 h showing very less decrease in the intensity or position of the absorption bands in case of complexes **1** and **2** indicating their excellent stability for about 12 h (a , b). Plot (c) shows gradual and rapid decay of curcumin (HCur) within the same time period indicating its instability in the buffer medium.



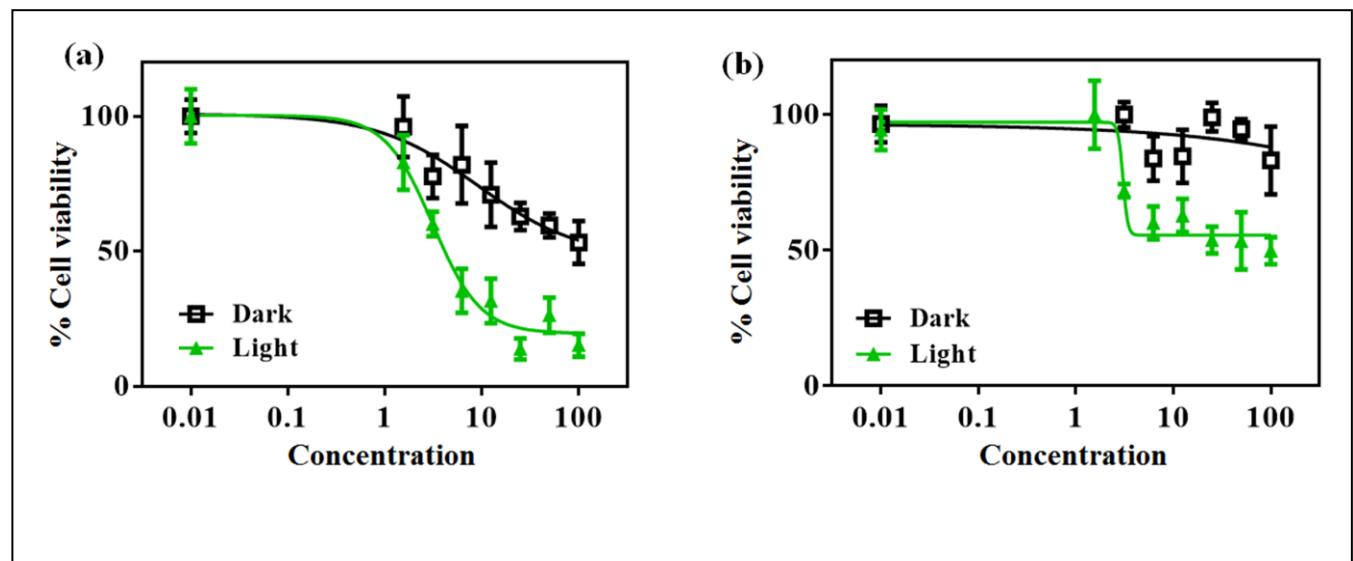
**Figure S11.** The UV-visible spectral traces of complex **2** in 2 % DMSO/DMEM (phenolate red free media; pH: 7.2) recorded over a period of 8 h showing very less decrease in the intensity or position of the absorption bands indicating its stability within the in vitro incubation time of 4 h.



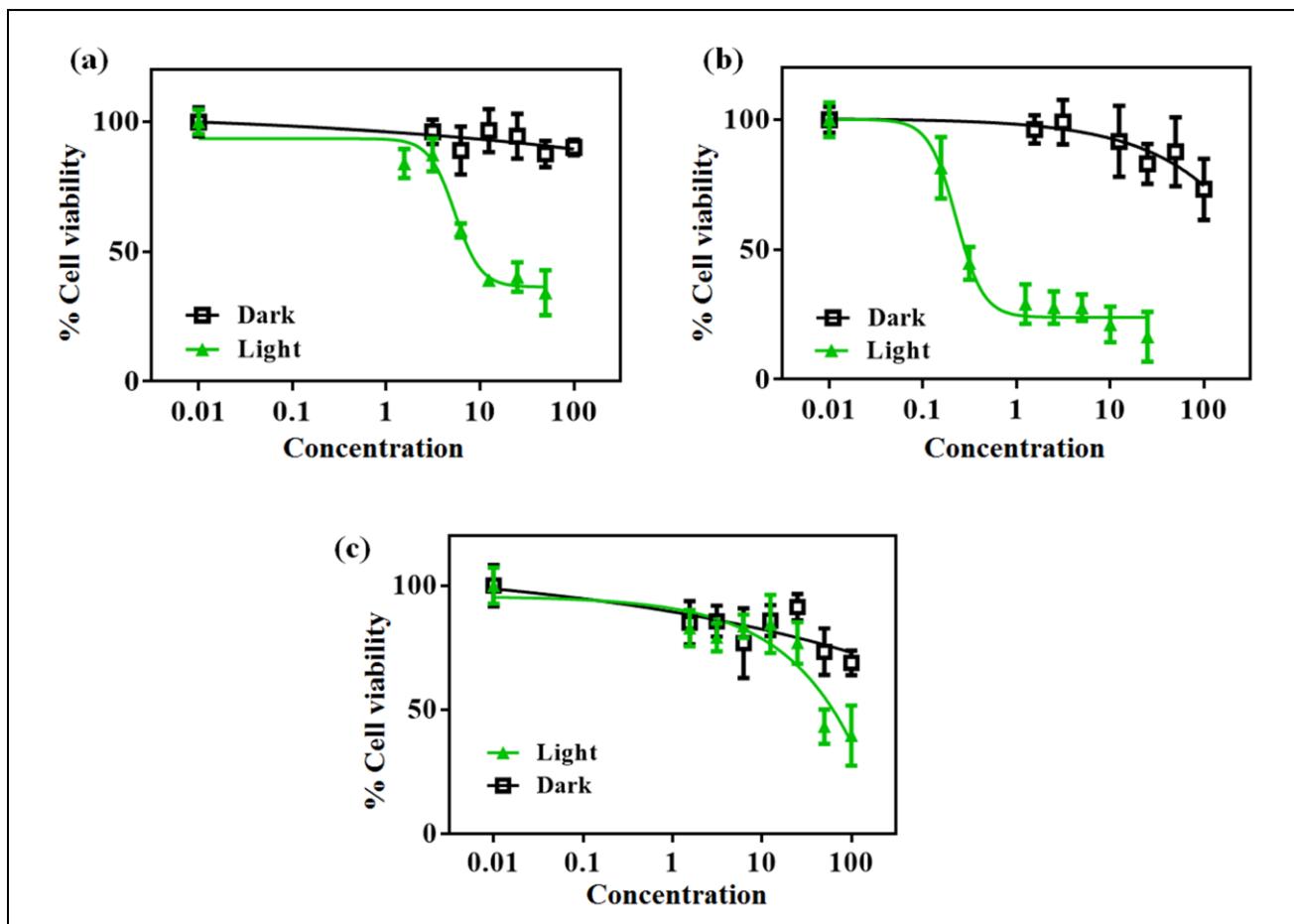
**Figure S12.** Photostability of the complexes and curcumin studied up to 48 h in 1:1 DMSO/DPBS medium upon 1 h irradiation with visible light (400-700 nm) with error bars. The instability of only curcumin is observed from the slope when compared to those of the complexes **1** and **2**.



**Figure S13.** Photocytotoxicity of complexes **1** and **2** in HeLa cells upon irradiation with visible light (400- 700 nm,  $10 \text{ J cm}^{-2}$ , green symbol) and in dark (black symbol) after 4 h incubation of the cells with the complexes: **1** (a), **2** (b).

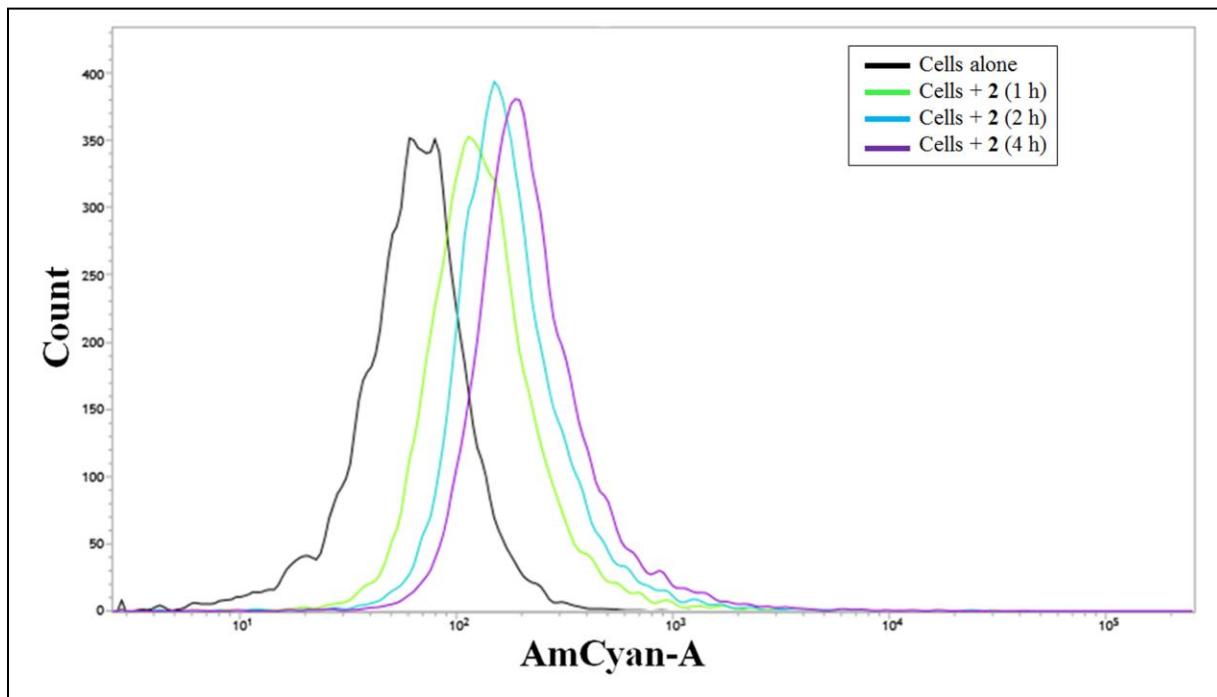


**Figure S14.** Photocytotoxicity of complexes **1** and **2** in MCF-7 cells upon irradiation with visible light (400- 700 nm,  $10 \text{ J cm}^{-2}$ , green symbol) and in dark (black symbol) after 4 h incubation of the cells with the complexes: **1** (a), **2** (b).

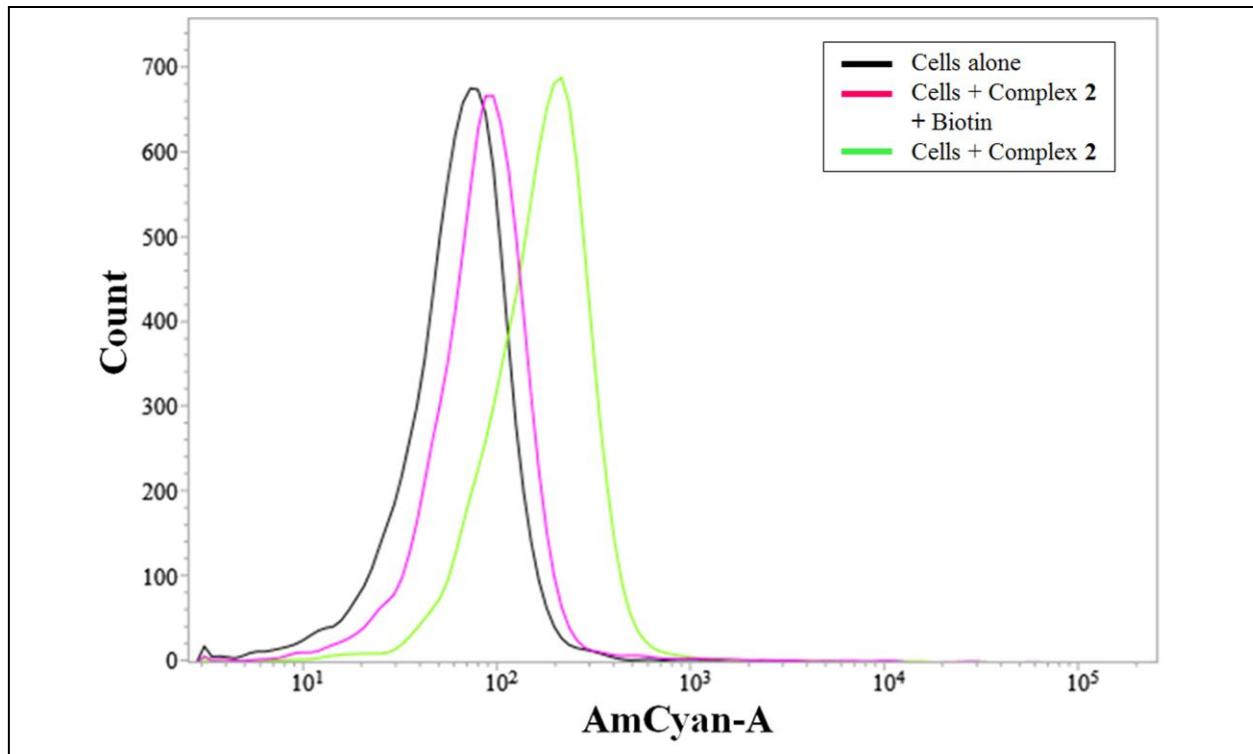


**Figure S15.** Photocytotoxicity of complexes **1** and **2** in HepG2 cells upon irradiation with visible light (400- 700 nm,  $10 \text{ J cm}^{-2}$ , green symbol) and in dark (black symbol) after 4 h incubation of the cells with the complexes: **1** (a), **2** (b).

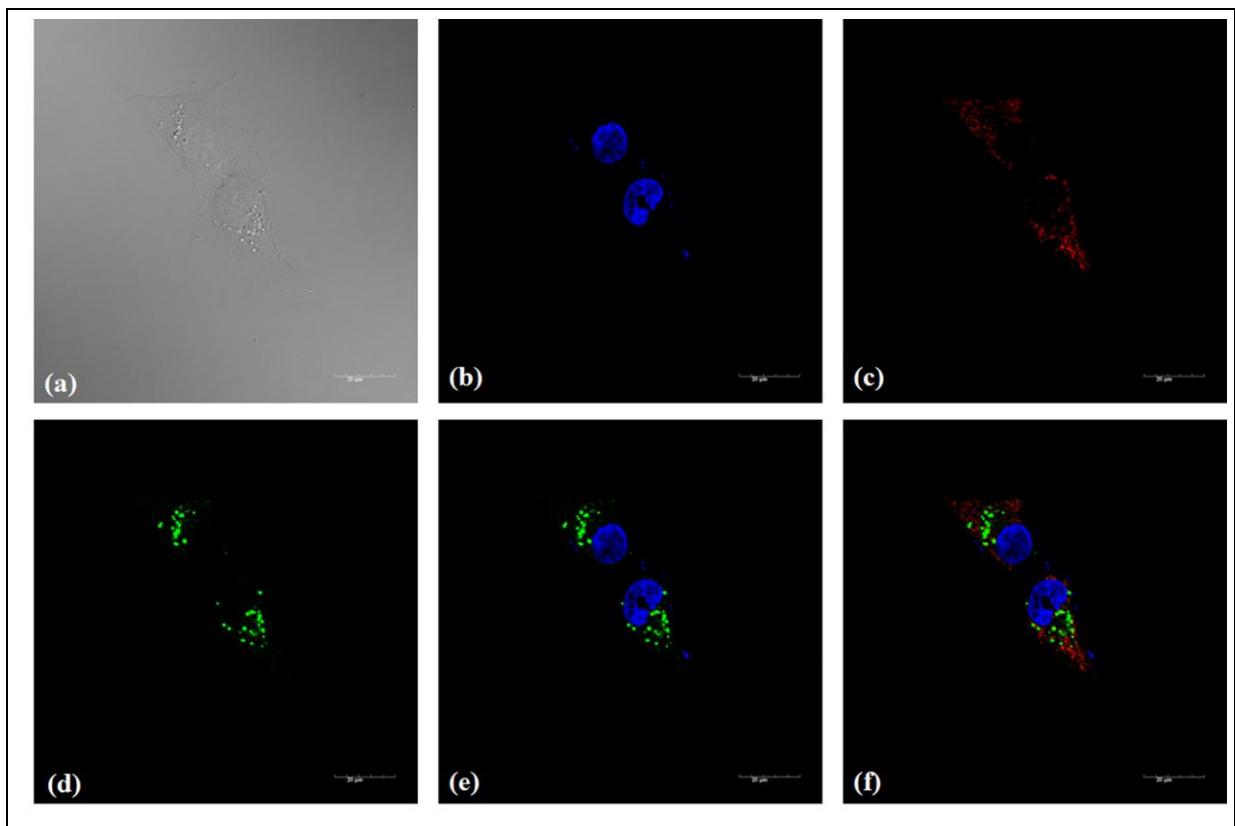
(c) An increase in the  $\text{IC}_{50}$  value upon saturation with 2 mM biotin (1 h) is shown in this plot. This corresponds to the histogram shown in Figure S14.



**Figure S16.** Histogram showing time dependent cellular incorporation of complex 2 ( $2 \mu\text{M}$ ) in HeLa cells at 1 h, 2 h and 4 h of incubation time at  $37^\circ\text{C}$ .

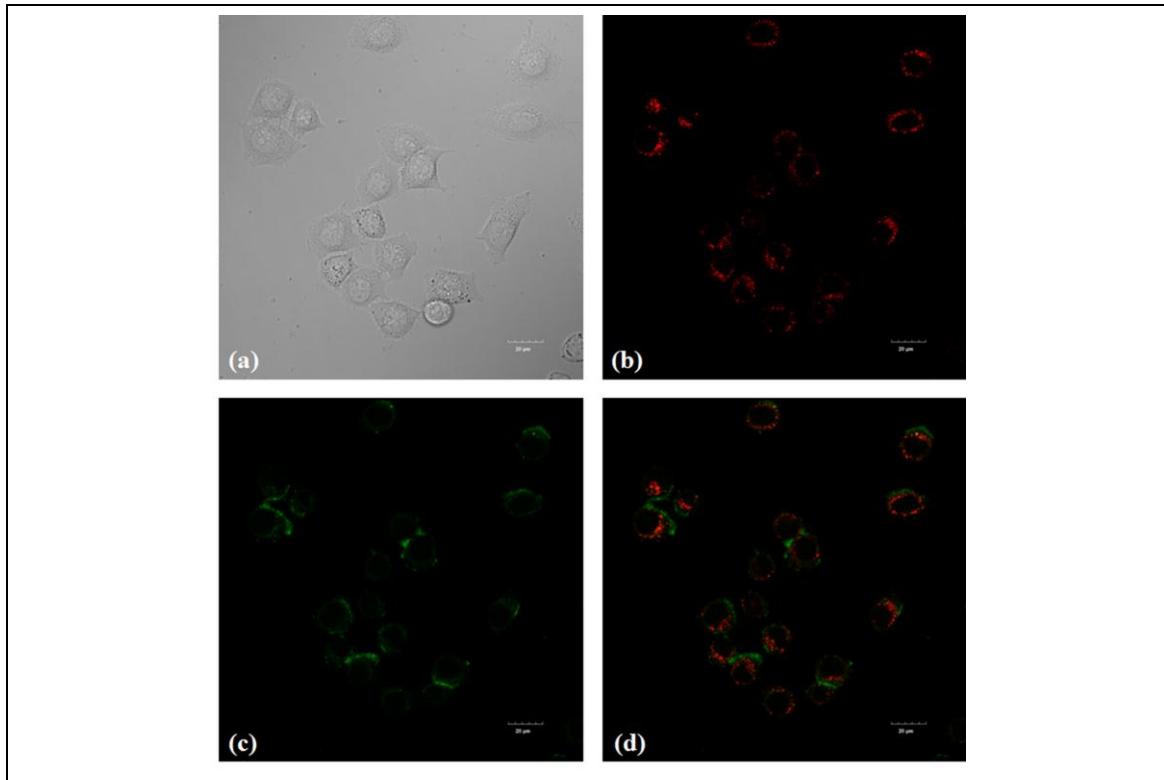


**Figure S17.** Histogram showing effect of pre-saturation with biotin (2 mM) on the cellular incorporation of the complex **2** (2  $\mu$ M) in HepG2 cells at 4 h incubation at 37°C. The uptake is lesser when cells were pre-saturated, as visible in Figure S15 (c) from MTT assay.

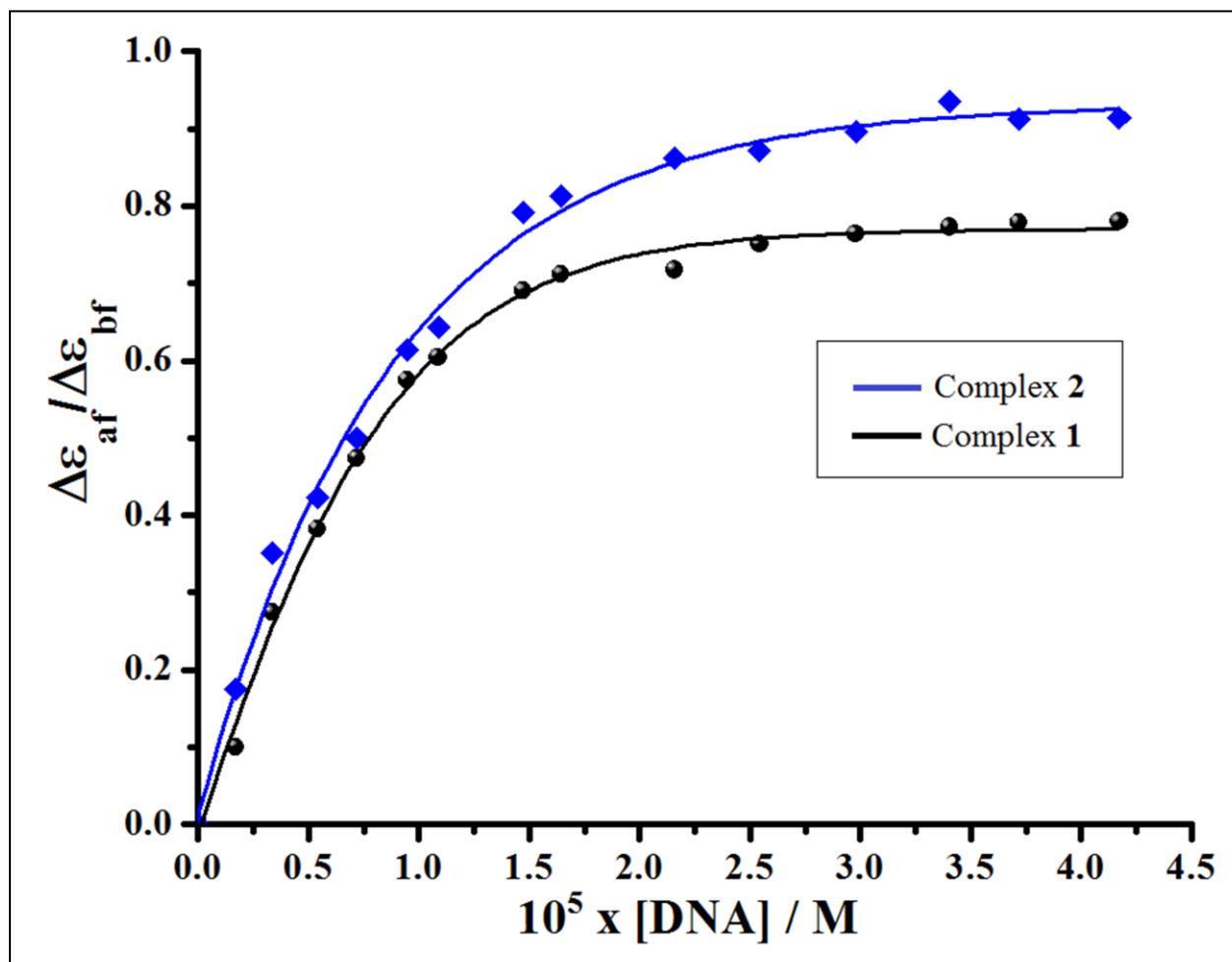


**Figure S18.** Confocal microscopy images of HeLa cells treated with complex **2** ( $20 \mu\text{M}$ ) co-stained with Mito tracker deep red: Panel (a) is bright field; panel (b) is for the blue fluorescence of Hoechst dye which localizes in the cell nucleus; panel (c) is for Mitotracker deep red; panel (d) is for the green fluorescence of complex **2**; panel (e) shows the merged image of (b) and (d) panels indicating non-nuclear localization of the complex; and panel (f) shows the merged image of the panels (b), (c) and (d) (Scale bar =  $20 \mu\text{m}$ ); Magnification: 100 X ; PCC: 0.35. This PCC value indicates only moderate localization of the complex in the mitochondria and non-nuclear localization.

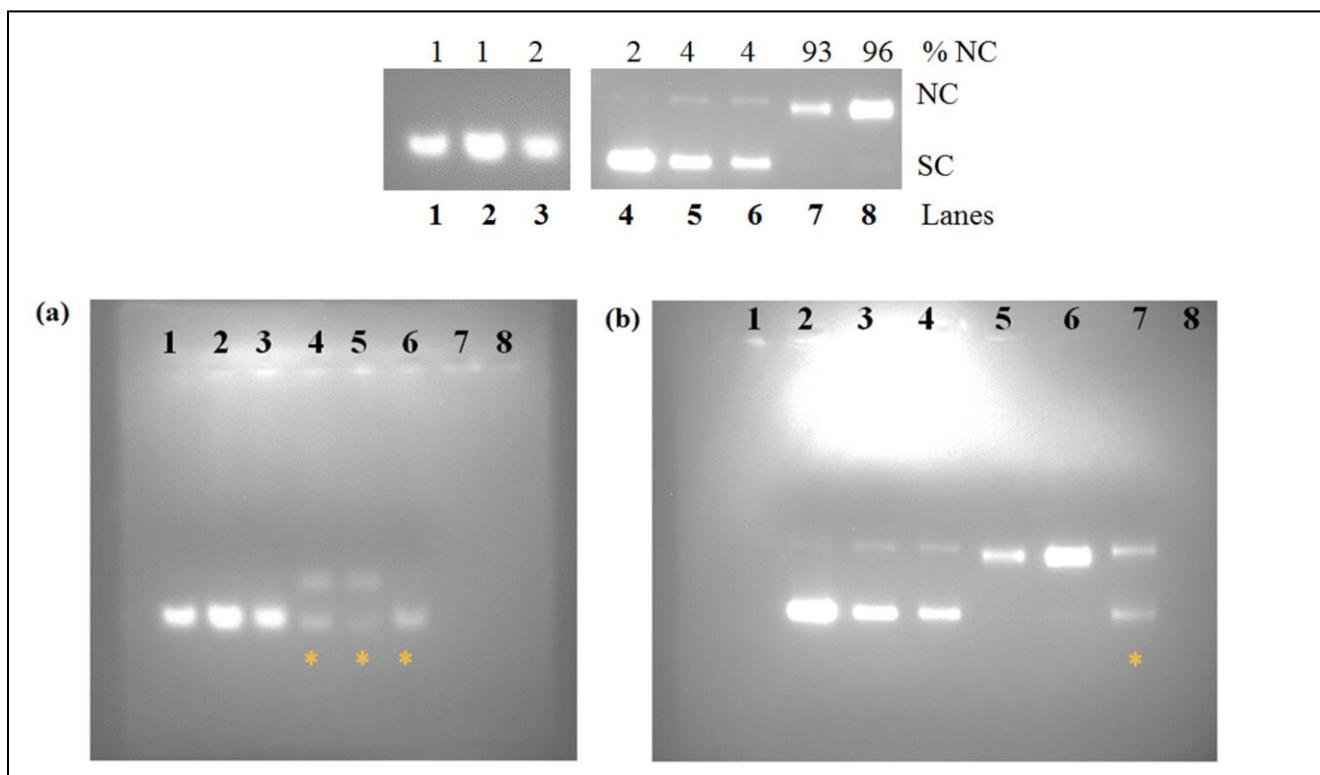
Panels (a) to (d) of this figure are the same and reproduced from Figure 9 of the main text for clarity and comparison.



**Figure S19.** Confocal microscopy images of HeLa cells treated with complex **2** (20  $\mu$ M) co-stained with Lysotracker deep red: Panel (a) is bright field; panel (b) is for Lysotracker deep red; panel (c) is for the green fluorescence of complex **2**; panel (d) shows the merged image of (b) and (c) (Scale bar = 20  $\mu$ m; Magnification: 60 X; PCC: 0.40). The PCC value indicates only moderate co-localization in the lysosomes.



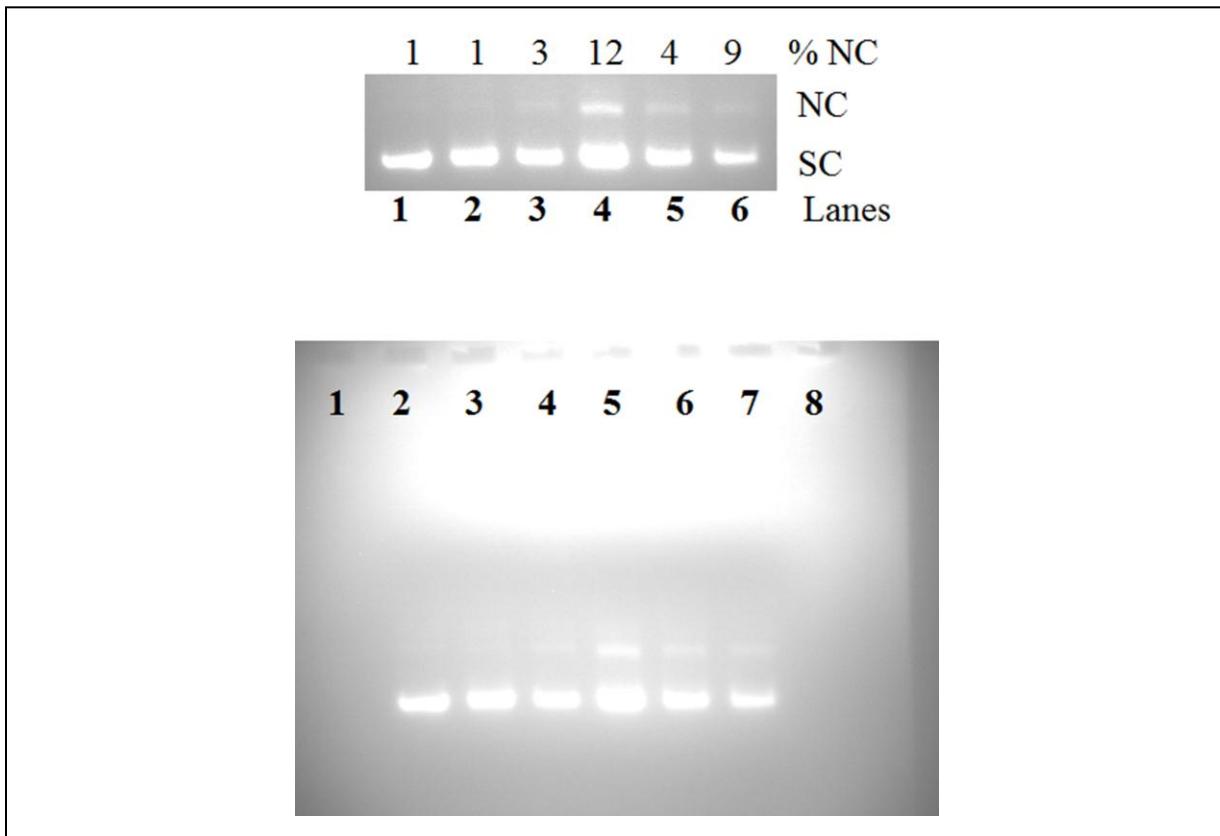
**Figure S20.** The plot of  $\Delta\epsilon_{af}/\Delta\epsilon_{bf}$  vs. [DNA] for the complexes **1** and **2** corresponding to their absorption spectral traces in 5 mM Tris-HCl buffer (pH 7.2) on increasing the quantity of calf thymus DNA.



**Figure S21.** Gel electrophoresis diagram showing the cleavage of pUC19 DNA (0.2 mg, 30 mM base pair) by monochromatic light of 446 nm diode laser, where [complex] = 20  $\mu$ M, incubation time: 1 h, exposure time: 1.5 h:

- (a) lane 1: DNA control (Dark); lanes 2 and 3: **1** (Dark) and **2** (Dark);
- (b) lane 2: DNA control (Light); lanes 3 and 4: ligand L<sup>1</sup> (Light) and ligand L<sup>2</sup> (Light); lanes 5 and 6: complex **1** (Light) and complex **2** (Light), where \* means the lanes that were not used for this study.

SC and NC are the supercoiled and the nicked circular form of plasmid DNA.

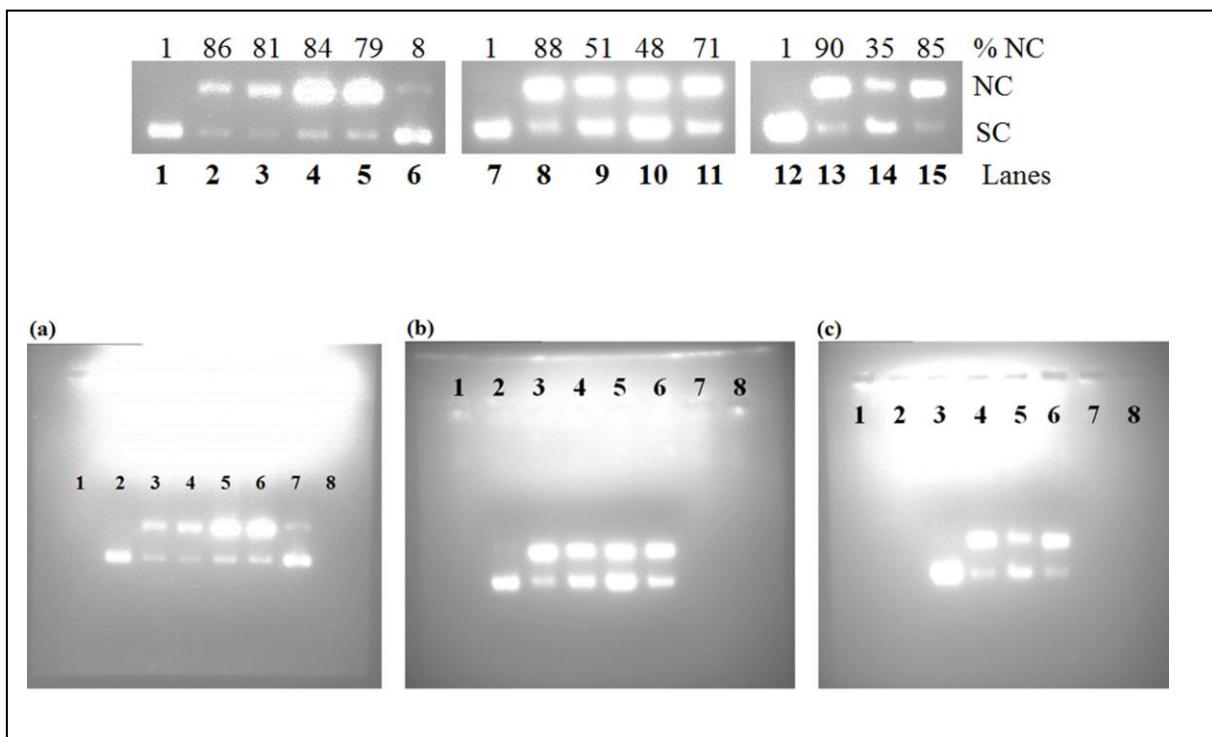


**Figure S22.** Gel electrophoresis diagram showing chemical nuclease activity of the complexes **1** and **2** (20  $\mu$ M) after incubating with GSH/H<sub>2</sub>O<sub>2</sub> (0.5 mM) for 1 h in dark (GSH is glutathione).

(Gel top): The lanes are taken from the raw gel that is shown below with lanes 1-8.

(Gel below): Lane 2: DNA control; lane 3: DNA + GSH; lane 4: DNA + **1** + GSH; lane 5: DNA + **1** + H<sub>2</sub>O<sub>2</sub>; lane 6: DNA + **2** + GSH; lane 7: DNA + **2** + H<sub>2</sub>O<sub>2</sub>.

SC and NC are the supercoiled and the nicked circular form of plasmid DNA.



**Figure S23.** Gel diagram showing the mechanistic aspects of photo nuclease activity of complex **2** in the presence of various singlet oxygen quenchers and hydroxyl radical scavengers in monochromatic diode laser light of 446 nm, where [complex] = 20  $\mu$ M, incubation time: 1 h, exposure time: 1 h. The details for the raw gels are given below (**the lanes shown in top are taken from the raw gels shown below**):

- (a) Lane 2: DNA control; lane 3: DNA + **2** (L); lane 4: DNA + **2** + TEMP (0.5 mM); lane 5: DNA + **2** + NaN<sub>3</sub> (0.5 mM); lane 6: DNA + **2** + DABCO (0.5 mM); lane 7: DNA + **2** under argon atmosphere (lanes are all light exposed).
- (b) Lane 2: DNA control; lane 3: DNA + **2** (L); lane 4: DNA + **2** + DMSO (4  $\mu$ L); lane 5: DNA + **2** + KI (0.5 mM); lane 6: DNA + **2** + SOD (superoxide dismutase, 4 units) (lanes are all light exposed).
- (c) Lane 3: DNA control; lane 4: DNA + **2** (L); lane 5: DNA + **2** + catalase (4 units); lane 6: DNA + **2** + D<sub>2</sub>O (16  $\mu$ L) (lanes are all light exposed).

**Table S1:** Computational data for the energy minimized structures of complex **1<sup>[S1]</sup>**

Charge 1

Multiplicity 6

Total Energy -2.163069845046529E+03

Dipole Moment -2.20056736E+00; -2.02004313E+00; -4.26429152E-04

<b>Atomic No</b>	<b>X</b>	<b>Y</b>	<b>Z</b>
7	-2.240844173	-6.930845519	-0.000115000
6	-2.627709202	-5.527734425	-0.000481000
6	-1.387281106	-4.579002349	-0.000119000
6	7.763659593	0.820029062	0.000051000
6	8.082501639	2.241568172	0.000033000
6	6.932873538	3.173463241	0.000043000
6	5.621884424	2.681147204	0.000031000
6	5.354362411	1.291732100	0.000028000
6	6.464026517	0.368107028	0.000048000
6	4.010838304	0.740783055	0.000019000
6	2.828645218	1.436783108	0.000019000
6	1.529309115	0.762243060	0.000024000
6	0.344234026	1.540147115	0.000090000
6	-0.965414071	1.008579075	0.000131000
6	-2.087149160	1.954472147	0.000138000
6	-3.402624258	1.574470118	0.000183000
6	-4.568825349	2.442195185	0.000169000

6	-4.481706343	3.853880295	-0.000101000
6	-5.627934454	4.659243358	-0.000102000
6	-6.973778516	4.046120309	0.000175000
6	-7.020235516	2.590418198	0.000441000
6	-5.874586457	1.827887139	0.000428000
8	9.302295733	2.653725202	0.000016000
8	7.047703519	4.534382345	0.000086000
8	-5.397384415	6.004739454	-0.000340000
8	-8.052988630	4.749082361	0.000186000
8	1.524755117	-0.558086043	-0.000045000
8	-1.221504093	-0.292799022	0.000173000
6	8.332433641	5.266312402	-0.000112000
6	-6.456945490	7.036633554	-0.000661000
7	-1.677588126	-3.083908236	-0.000034000
6	-2.357824180	-2.641081199	1.274433095
6	-1.317005103	-2.461965189	2.365699179
6	-1.574994120	-2.642217200	3.730916282
6	-0.548290041	-2.389582181	4.661647357
6	0.716995055	-1.974729153	4.205229323
6	0.923581070	-1.825127141	2.827326217
7	-0.078797006	-2.058002157	1.946578151
6	-2.357810178	-2.640881202	-1.274443096
6	-1.316985100	-2.461873186	-2.365716182
6	-1.574936122	-2.642212204	-3.730927284
6	-0.548194043	-2.389661182	-4.661642356

6	0.717082052	-1.974796152	-4.205212321
6	0.923632072	-1.825108138	-2.827313215
7	-0.078777006	-2.057905158	-1.946580151
26	0.088252007	-1.916211145	0.000006000
17	1.732475131	-3.651174277	-0.000007000
1	-3.523342268	4.362828331	-0.000334000
1	-8.011092609	2.145303163	0.000647000
1	-5.942561444	0.741928056	0.000628000
1	-3.605267277	0.503583038	0.000254000
1	-1.813710136	3.006916227	0.000113000
1	0.452530035	2.619414201	0.000108000
1	2.806884216	2.523748194	0.000023000
1	3.933163302	-0.346650026	0.000018000
1	8.610939680	0.140452011	0.000062000
1	6.255130464	-0.699270052	0.000059000
1	4.820109368	3.412509263	0.000033000
1	8.031014598	6.315359461	-0.000190000
1	8.914511681	5.018723385	0.890706065
1	8.914342662	5.018550381	-0.890994070
1	-5.901240427	7.975955610	-0.001019000
1	-7.082240537	6.942810534	-0.891601068
1	-7.082159520	6.943434540	0.890399069
1	1.876631142	-1.535184115	-2.401819183
1	1.530730115	-1.781581134	-4.895361373
1	-0.731039055	-2.522768194	-5.723871432

1	-2.554836193	-2.973065230	-4.060449312
1	-2.823347217	-1.669083126	-1.084005085
1	-3.133668239	-3.345258253	-1.591063121
1	-3.257042247	-5.344174426	-0.880989069
1	-3.257830250	-5.343864434	0.879405067
1	-0.764246059	-4.792183367	-0.873295065
1	-0.764658061	-4.792318365	0.873323069
1	-3.133519238	-3.345646257	1.591040123
1	-2.823581213	-1.669368130	1.084105083
1	1.530619115	-1.781452137	4.895389372
1	-0.731169057	-2.522617194	5.723879422
1	-2.554897196	-2.973071225	4.060430309
1	1.876591141	-1.535218115	2.401846182
1	-1.877039143	-7.331405550	-0.854890063
1	-1.877310145	-7.331043540	0.854947066

**Table S2:** Computational data for the energy minimized structures of complex **2**

Charge 1

Multiplicity 6

Total Energy -2.823882612948552E+03

Dipole Moment 2.20387029E+00; -7.94783954E-01; 7.94519377E-01

<b>Atomic No</b>	<b>X</b>	<b>Y</b>	<b>Z</b>
7	-6.471773460	2.291346564	-2.316057247
6	-7.921097498	2.063496006	-2.159107857
6	-8.091835214	0.630281304	-2.800945397
7	-6.860208876	0.506610102	-3.590950218
6	-5.872618076	1.406306034	-3.205698676
6	-8.411376674	2.071389965	-0.690313732
6	-8.258042486	-0.438691483	-1.667683436
16	-9.366939444	0.451555543	-0.399384415
8	-4.668571234	1.401931812	-3.568424947
6	-6.921416850	-0.913241444	-1.072150149
6	-6.997353468	-2.005950975	0.014635071
6	-5.560283905	-2.340312398	0.466882058
6	-5.426765130	-3.467191836	1.524220039
6	-3.941876195	-3.689177760	1.783346178
7	-3.442583348	-3.245412683	2.988209796
8	-3.175259226	-4.174842341	0.900489886
6	-1.997653333	-3.214697526	3.254288149
6	-1.336773810	-1.872274838	2.788393123
6	2.148596894	7.994138246	0.734664866
6	3.215933176	8.471522748	-0.036135148
6	4.080995629	7.552767537	-0.700520085
6	3.877311560	6.179886434	-0.591677040
6	2.793107795	5.682250636	0.191318074

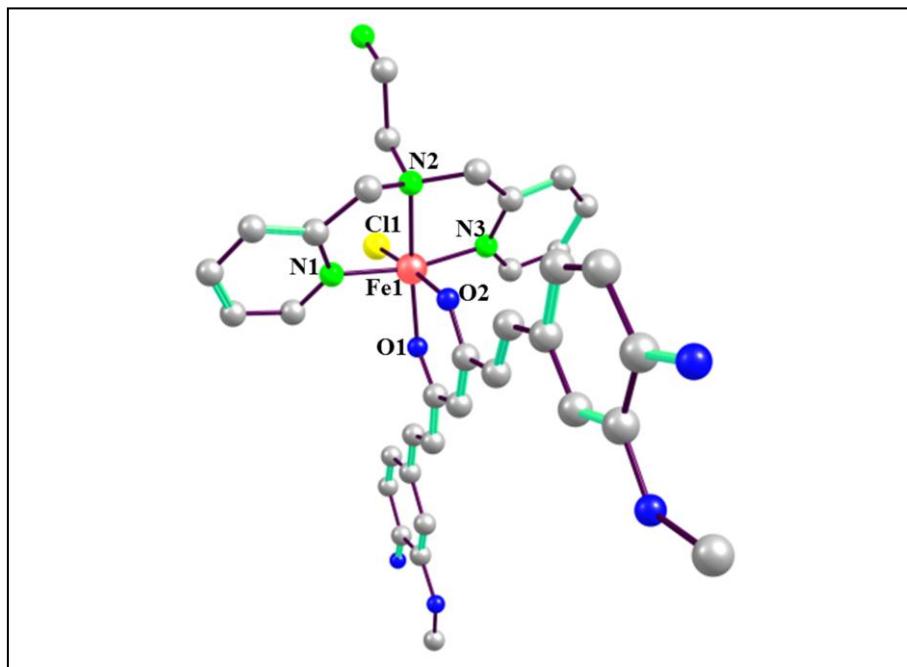
6	1.942558663	6.611819849	0.844430393
6	2.515426548	4.263961531	0.352522203
6	3.200713567	3.201450029	-0.178614133
6	2.793741824	1.827783848	0.072785124
6	3.513986663	0.749807040	-0.484734727
6	3.185393018	-0.616736520	-0.295619376
6	4.030647542	-1.625692867	-0.932282141
6	3.802500348	-2.970280598	-0.833483838
6	4.573982393	-4.059551997	-1.418195927
6	5.740627089	-3.845072431	-2.210002246
6	6.432039075	-4.933106353	-2.737165960
6	5.986539355	-6.265463444	-2.494207900
6	4.842102974	-6.487556119	-1.719045031
6	4.146077123	-5.391900022	-1.188243081
8	3.430792151	9.828219600	-0.153570997
8	5.090462341	8.194389183	-1.424377765
8	7.584214054	-4.888294830	-3.526539631
8	6.681505084	-7.333064097	-3.021867797
8	1.721933026	1.633520854	0.844506288
8	2.148675408	-1.011597063	0.433156796
7	-0.112543077	-2.007457882	1.917770567
6	-0.433229028	-2.584803601	0.564377114
6	-1.063826768	-1.525962382	-0.320287691
6	-2.012142344	-1.831806082	-1.309823855
6	-2.515162520	-0.793395666	-2.114899707

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6	2.937643951	-1.280620521	5.622780479
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6	2.321640934	0.455541162	4.056956880
7	1.710640384	-0.457265392	3.261695292
26	0.732315610	0.077457451	1.438999382
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1	4.518560602	-7.508734645	-1.547571573
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1	1.113487761	6.241268772	1.442361785
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6	6.058280973	7.391471983	-2.163513575
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1	6.625511093	6.743496558	-1.482393753
1	5.556295020	6.787913739	-2.931106040
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1	4.376106864	0.982514196	-1.100794122
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1	2.935588262	-3.272755705	-0.246211084
1	1.654303375	4.012702072	0.972548222
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1	3.413358164	-1.603145635	6.544461406
1	2.249879605	-3.266758089	5.065744906
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1	-6.305776854	-1.303843353	-1.896630385
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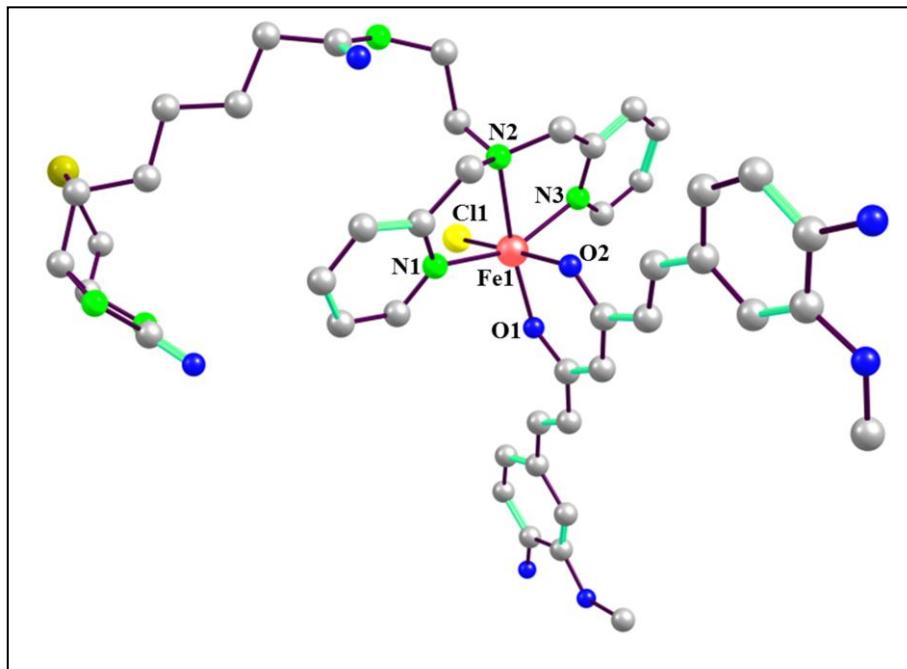
1	-1.106918994	-3.446640005	0.634545472
1	0.512285914	-2.896050233	0.107901714
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1	0.688245999	-3.626825752	3.115642826
1	-8.491525066	2.796829690	-2.744009836
1	-8.980929406	0.604304051	-3.437458462
1	-4.970024208	-2.639048624	-0.409757077
1	-5.095315288	-1.423093764	0.861889882
1	2.280111914	1.487572430	3.727517155
1	-0.771171291	1.762959958	-0.666087363
1	-2.487084746	1.335889793	-2.500893033
1	-3.254947028	-0.986131720	-2.885428164
1	-2.357689251	-2.854973717	-1.416534686
1	-8.846677402	-1.290673788	-2.028772374
1	-7.567545288	2.116078317	0.002651627

**Table S3: Estimated Bond Distances [Å] and Angles [°] for the Complex 1 from the DFT Calculations**



Bond	Bond Length (Å)	Bond	Bond Angle (°)
Fe(1)-O(1)	1.9769	N(1)-Fe(1)-O(1)	96.41
Fe(1)-O(2)	2.0858	O(1)-Fe(1)-O(2)	85.50
Fe(1)-N(1)	1.9588	Cl(1)-Fe(1)-O(2)	175.43
Fe(1)-N(2)	2.117	Cl(1)-Fe(1)-N(2)	99.98
Fe(1)-N(3)	1.9588	N(1)-Fe(1)-N(3)	167.16
Fe(1)-Cl(1)	2.3903	O(1)-Fe(1)-N(2)	170.08
		N(3)-Fe(1)-O(2)	90.16

**Table S4: Estimated Bond Distances [Å] and Angles [°] for the Complex 2 from the DFT Calculations**



Bond	Bond Length (Å)	Bond	Bond Angle (°)
Fe(1)-O(1)	1.9376	N(1)-Fe(1)-O(1)	103.13
Fe(1)-O(2)	2.0503	O(1)-Fe(1)-O(2)	85.6
Fe(1)-N(1)	2.1116	Cl(1)-Fe(1)-O(2)	176.45
Fe(1)-N(2)	2.2999	Cl(1)-Fe(1)-N(2)	100.13
Fe(1)-N(3)	2.1366	N(1)-Fe(1)-N(3)	153.39
Fe(1)-Cl(1)	2.3637	O(1)-Fe(1)-N(2)	168.38
		N(3)-Fe(1)-O(2)	88.24

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