

Supplementary Information

Ferrocene functionalized dithiocarbamate complexes holding thiophene arms: Synthesis, crystal structure, electrochemical, *in silico* interaction profiling, and apoptosis induction in HeLa Cells

Bhaveshkumar B. Makwana^a, Heena A. Parmar^b, Adam N. Khan^c, José M. Padrón^c, and Umesh P. Tarpada^{d*}

^a Department of Chemistry, School of Sciences, Gujarat University, Ahmedabad- 380009, Gujarat, India.

^b Department of Chemistry, Gujarat Arts and Science College, Ahmedabad- 380006, Gujarat, India.

^c BioLab, Instituto Universitario de Bio-Orgánica “Antonio González”, Universidad de la Laguna, C/ Astrofísico Francisco Sánchez 2, 38206 La Laguna, Spain.

^d Department of Chemistry, Government Arts and Science College, Bavla, Ahmedabad- 382220, Gujarat, India.

*Corresponding author: umeshtarpada@gmail.com (U.P. Tarpada)

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General procedure for the syntheses of L¹/L² ligands

An equivalent amount of *formylferrocene* (0.137 g, 1 mmol) was added to a solution containing thiophenemethylamine (0.122 g, 1 mmol) or thiopheneethylamine (0.172 g, 1 mmol) in absolute alcohol (10 mL) at room temperature. After five hours of stirring of the reaction mixture, thin layer chromatography (TLC) was used to verify that the completion of reaction. The reaction mixture was filtered through a celite pad, dried over Na₂SO₄, and the solvent was extracted under vacuum to reveal the ligands L¹ and L².

N-thiophenemethyl-*N*-methylferrocenylamine (L¹)

M.W. 311.22; Reddish brown solid; Yield: *ca* 281.0 mg, 90.3%; m.p. 159-163 °C dec. ¹H NMR (400 MHz, DMSO-*d*₆) δ ppm: 7.39-7.37 (m, 1H, ThH-5); 6.98-6.96 (m, 2H, ThH-3 & ThH-4); 4.20-4.18 (m, 2H, Fc); 4.12 (s, 7H, Fc); 4.09-4.08 (m, 2H, -CH₂Th); 3.87 (s, 2H, -CH₂Fc). ¹³C{¹H} NMR (400 MHz, DMSO-*d*₆) δ ppm: 144.7, 127.1, 125.9, 125.6, 125.3, 125.2, 118.5, 86.5, 70.2, 68.8, 68.1, 67.8, 47.4 (-CH₂Fc), 47.2 (-CH₂Th). Infrared spectrum (KBr disc, cm⁻¹): 3922 w, 3321 w (ν(N-H)), 3086 s (aromatic, ν(C-H)), 2813 s (arylidene, ν(C-H)), 1634 m, 1438 s (ν(C=N)), 1328 w, 1227 m, 1101 s, 1038 m (ν(C-C)), 997 m, 923 w, 813 s, 697 s (aromatic out-of-plane bending, γ(C-H))/(thiophene ν(C-S)), 611 m, 482 s (ν(Fe-Cp)), 415 w. UV-visible (DCM, λ, nm): 306, 272.

N-thiopheneethyl-*N*-methylferrocenylamine (L²)

M.W. 325.25; Reddish brown solid; Yield: *ca* 291.7 mg, 89.7%; m.p. 163-168 °C dec. ¹H NMR (400 MHz, DMSO-*d*₆) δ ppm: 7.33 (m, 1H, ThH-5); 6.96-6.91 (m, 2H, ThH-3 & ThH-4); 4.21 (s, 2H, Fc); 4.09 (s, 7H, Fc); 3.48 (s, 2H, -CH₂CH₂Th); 2.98 (s, 2H, -CH₂Fc); 2.84 (s, 2H, -CH₂CH₂Th). ¹³C{¹H} NMR (400 MHz, DMSO-*d*₆) δ ppm: 143.4, 142.9, 127.3, 127.0, 125.5, 125.1, 124.3, 124.1, 87.0, 83.3, 73.1, 70.2, 69.3, 68.7, 68.6, 68.0, 67.9, 67.7, 59.6, 52.6, 50.8 (-CH₂Fc), 50.2, 48.2 (-CH₂Th), 30.0, 27.6. Infrared spectrum (KBr disc, cm⁻¹):

3813 w, 3640 w ($\nu(\text{N-H})$), 3086 s (aromatic, $\nu(\text{C-H})$), 2915 w, 2815 s (arylidene, $\nu(\text{C-H})$),
2250 w, 1641 m, 1438 s ($\nu(\text{C=N})$), 1321 w, 1229 m, 1104 s ($\nu(\text{C-C})$), 998 m, 920 m, 814 s,
692 s (aromatic out-of-plane bending, $\gamma(\text{C-H})$)/(thiophene $\nu(\text{C-S})$), 480 s ($\nu(\text{Fe-Cp})$), 409 w.
UV-visible (DCM, λ , nm): 318, 270.

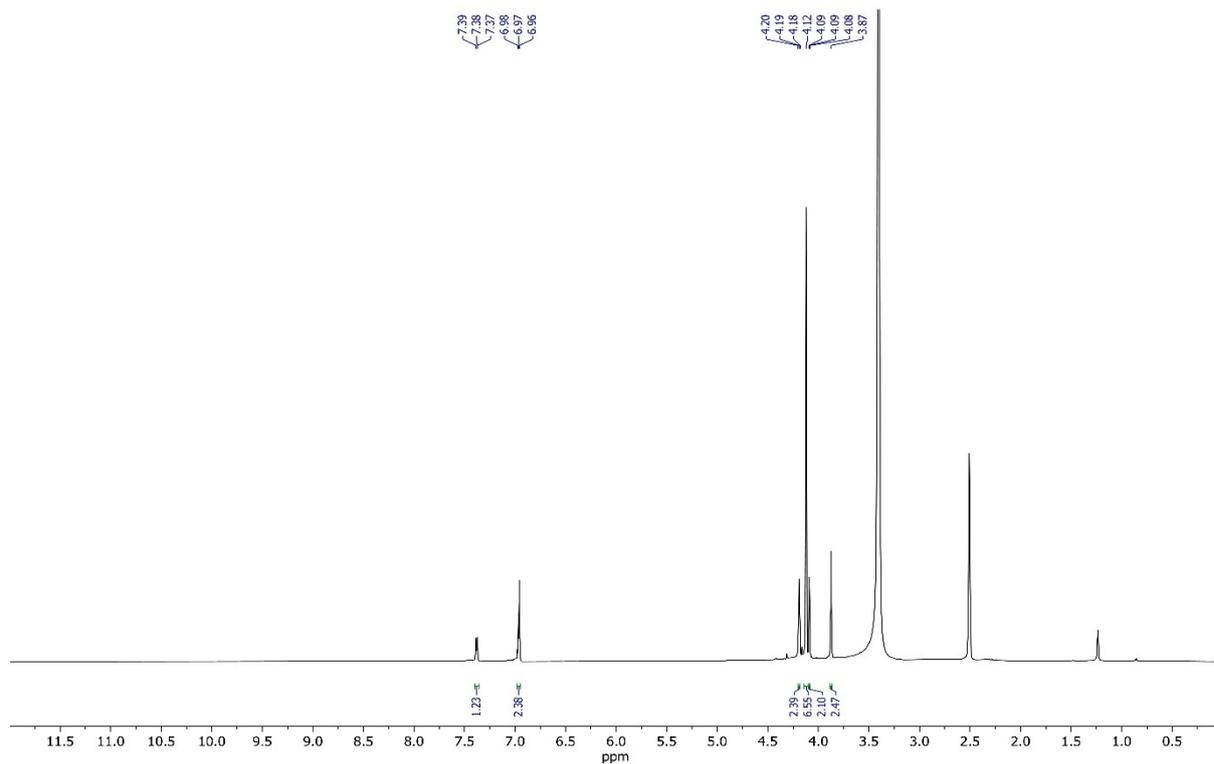


Fig. S1. ^1H NMR spectra (400 MHz, $\text{DMSO-}d_6$) of ligand L^1 .

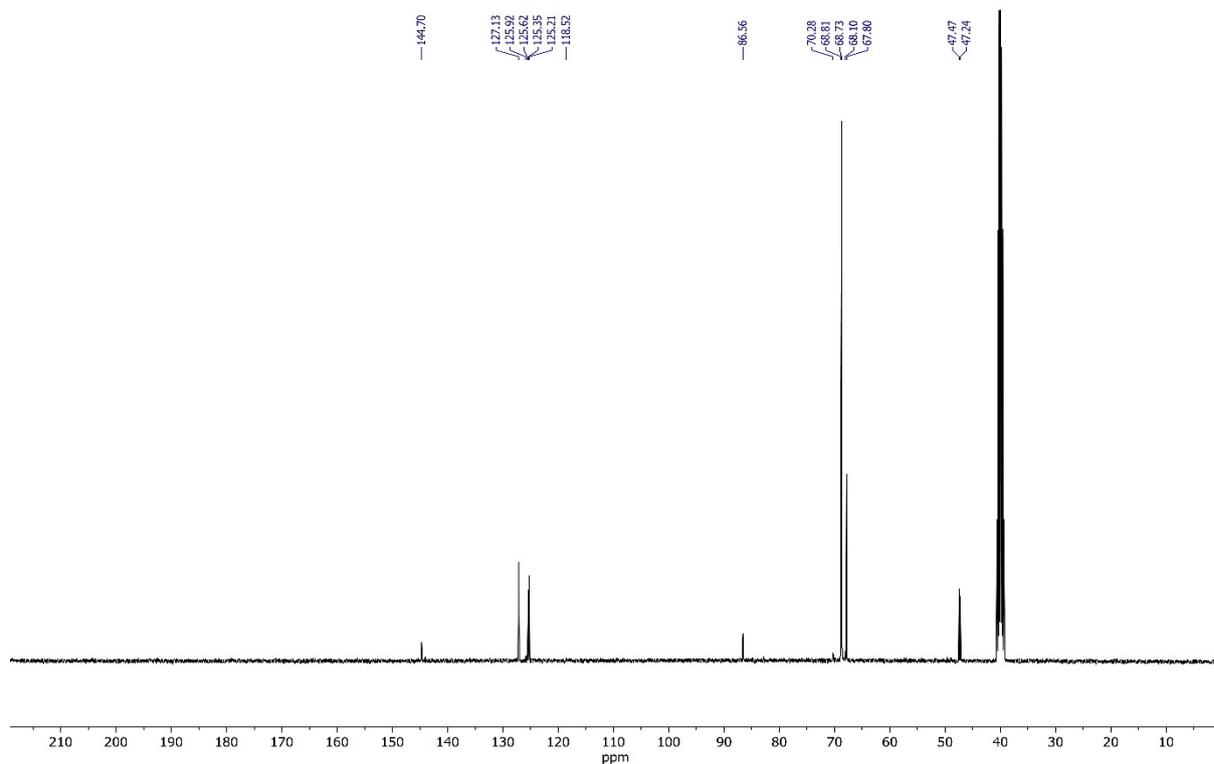


Fig. S2. $^{13}\text{C}\{^1\text{H}\}$ NMR spectra (400 MHz, $\text{DMSO-}d_6$) of ligand L^1 .

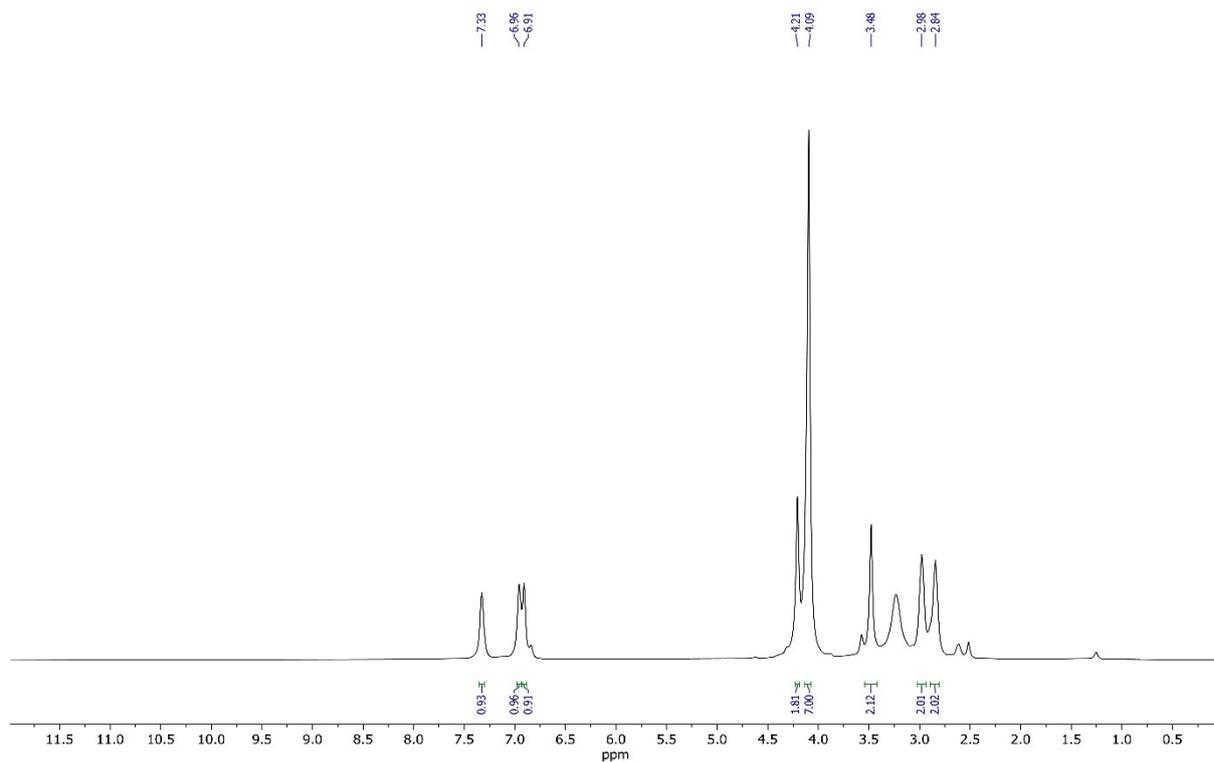


Fig. S3. ^1H NMR spectra (400 MHz, $\text{DMSO-}d_6$) of ligand L^2 .

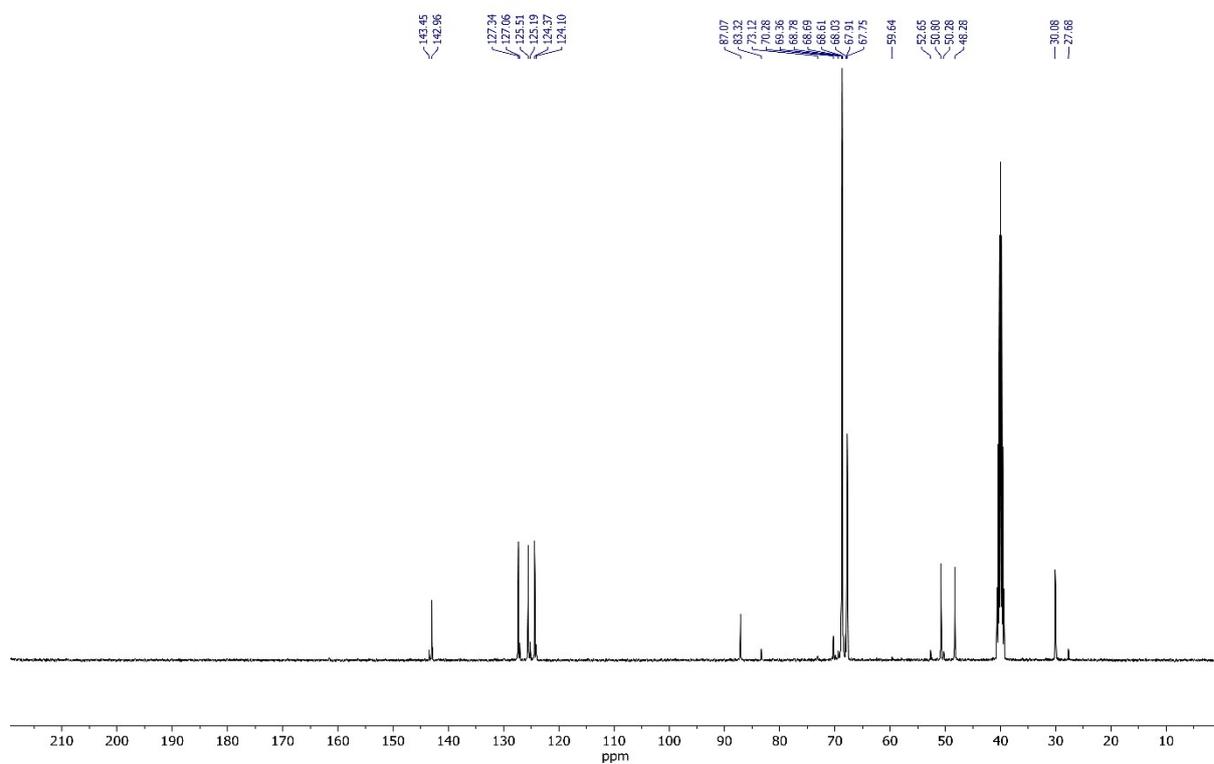


Fig. S4. $^{13}\text{C}\{^1\text{H}\}$ NMR spectra (400 MHz, $\text{DMSO-}d_6$) of ligand L^2 .

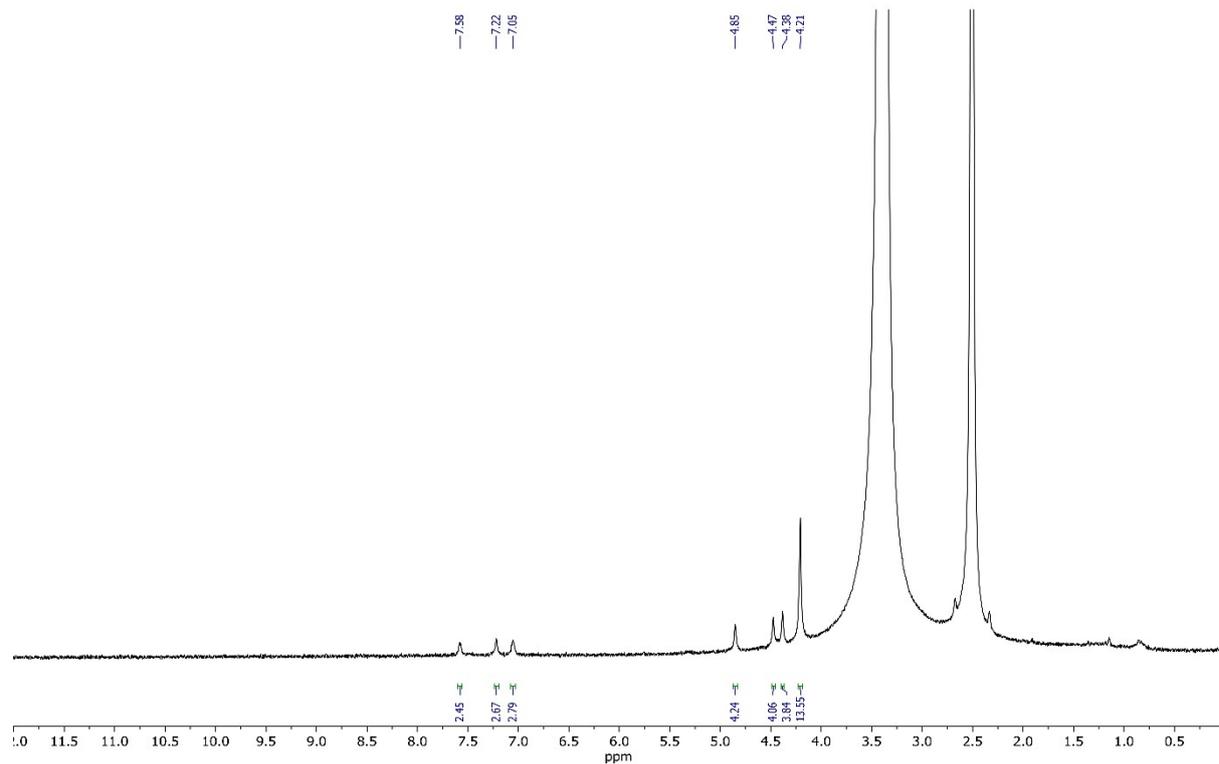


Fig. S5. ^1H NMR spectra (400 MHz, $\text{DMSO-}d_6$) of complex **1**.

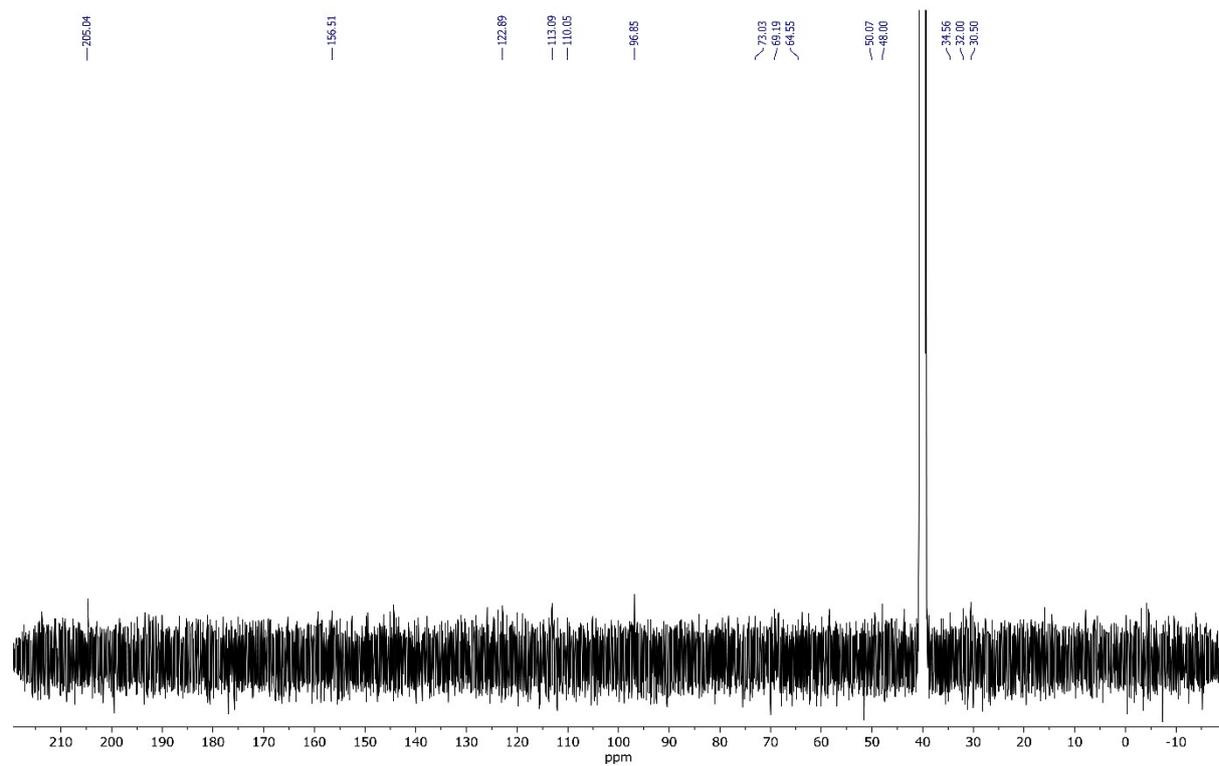


Fig. S6. $^{13}\text{C}\{^1\text{H}\}$ NMR spectra (400 MHz, $\text{DMSO-}d_6$) of complex **1**.

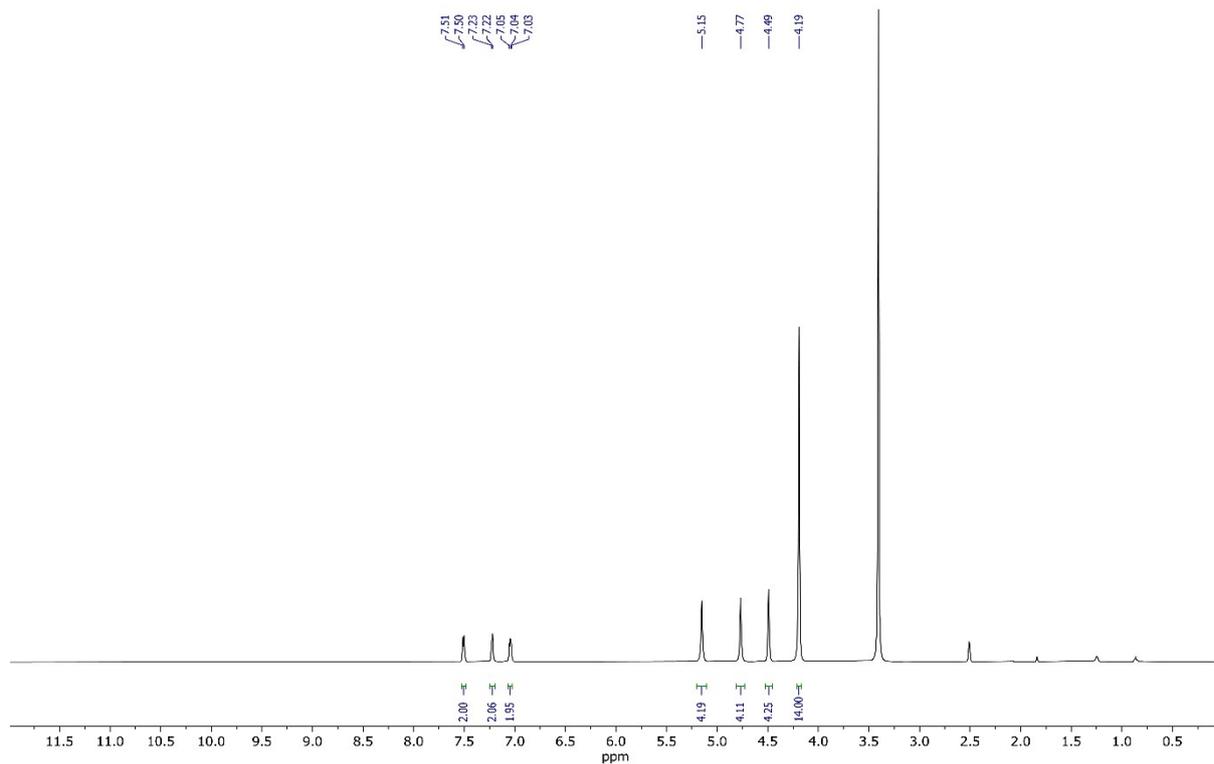


Fig. S7. ^1H NMR spectra (400 MHz, $\text{DMSO-}d_6$) of complex **3**.

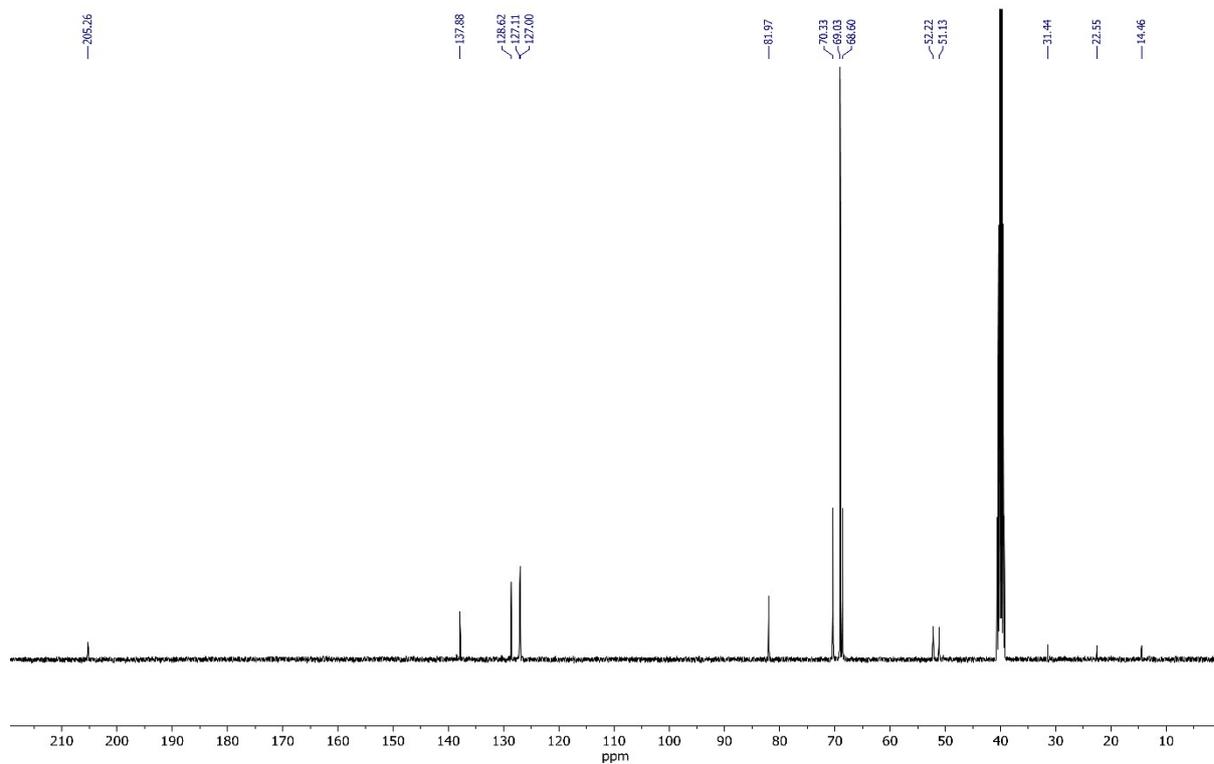


Fig. S8. $^{13}\text{C}\{^1\text{H}\}$ NMR spectra (400 MHz, $\text{DMSO-}d_6$) of complex **3**.

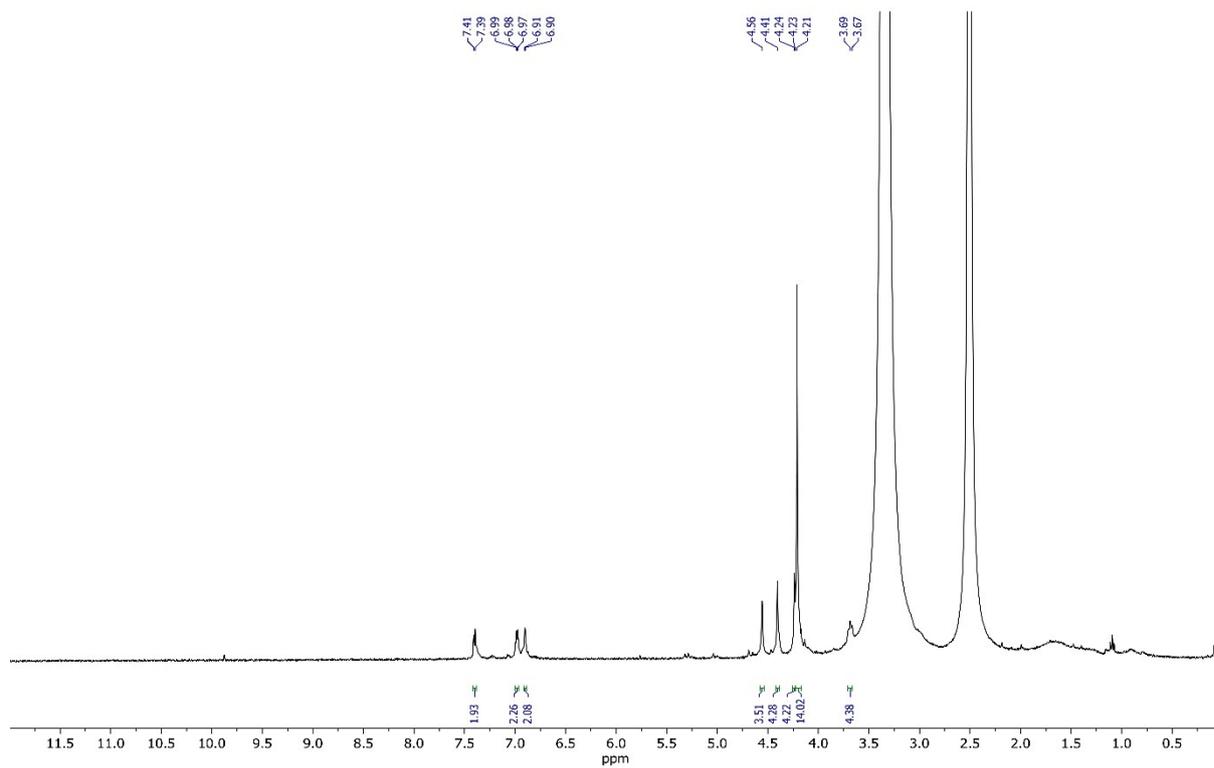


Fig. S9. ^1H NMR spectra (400 MHz, $\text{DMSO-}d_6$) of complex **4**.

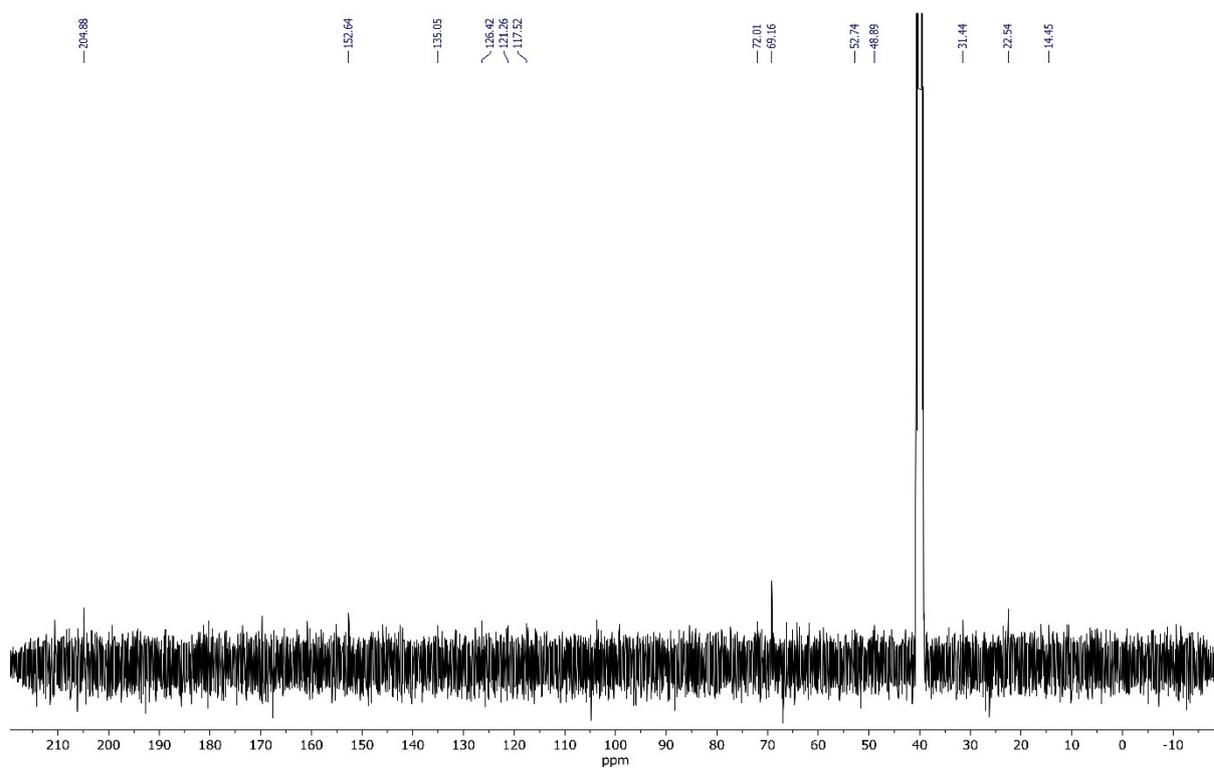


Fig. S10. $^{13}\text{C}\{^1\text{H}\}$ NMR spectra (400 MHz, $\text{DMSO-}d_6$) of complex **4**.

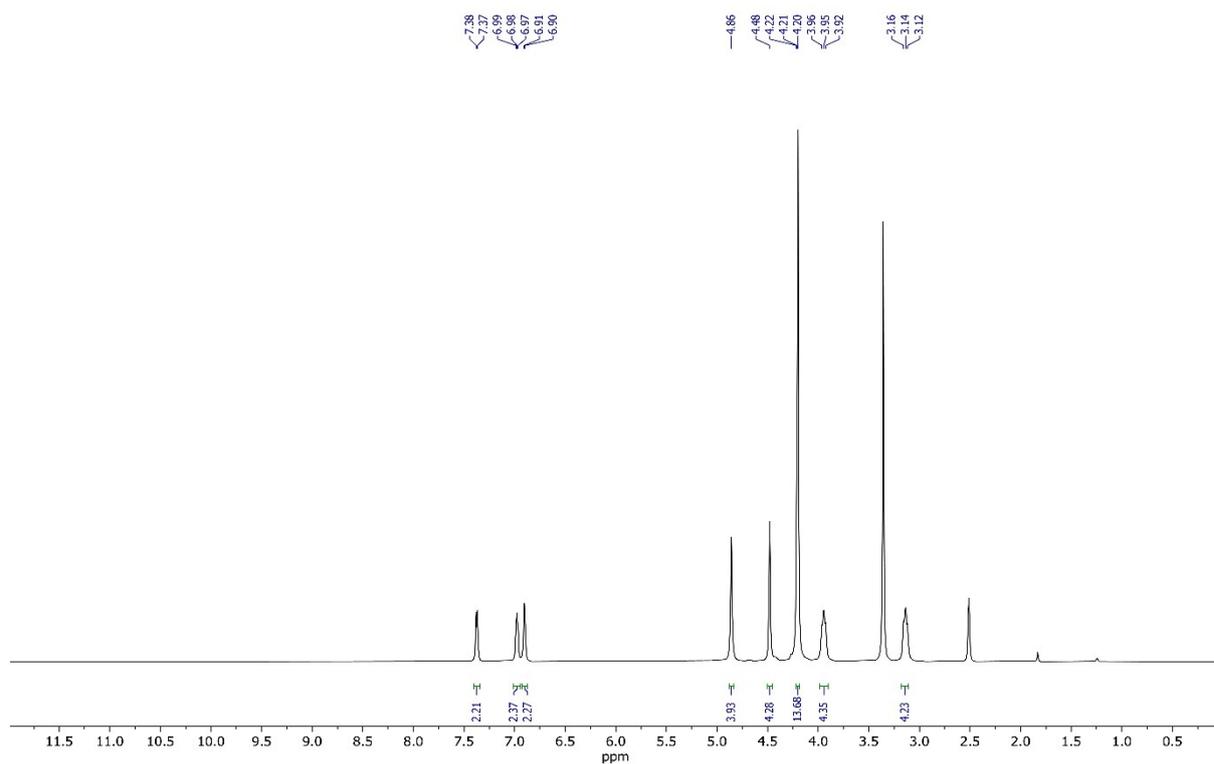


Fig. S11. ^1H NMR spectra (400 MHz, $\text{DMSO-}d_6$) of complex **6**.

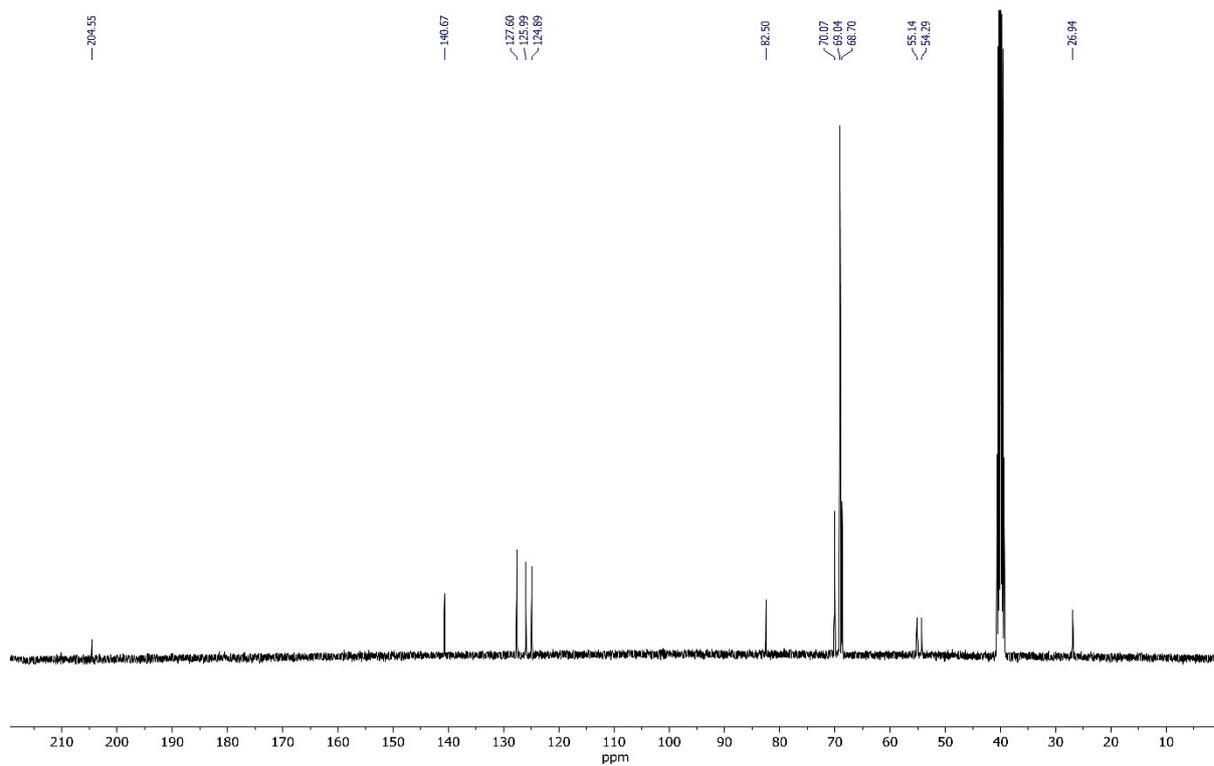


Fig. S12. $^{13}\text{C}\{^1\text{H}\}$ NMR spectra (400 MHz, $\text{DMSO-}d_6$) of complex **6**.

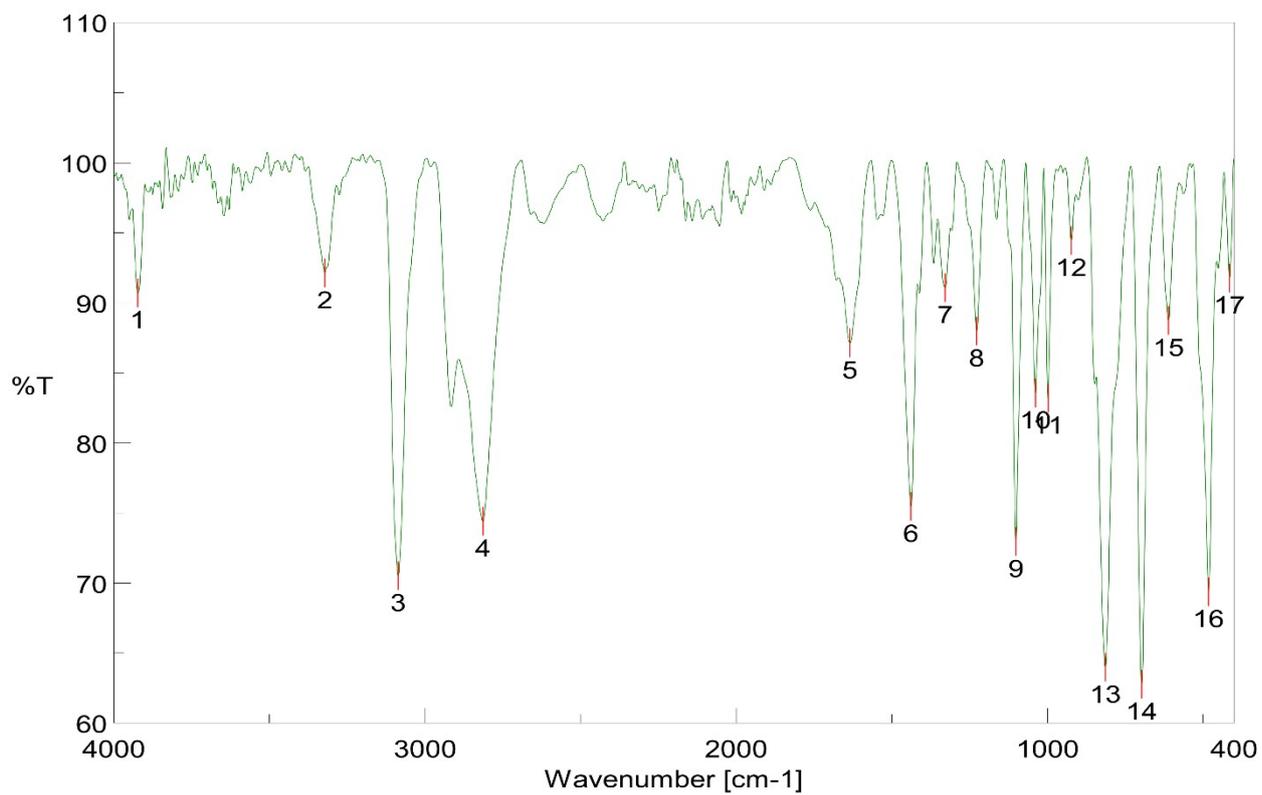


Fig. S13. FT-IR spectra of ligand L¹.

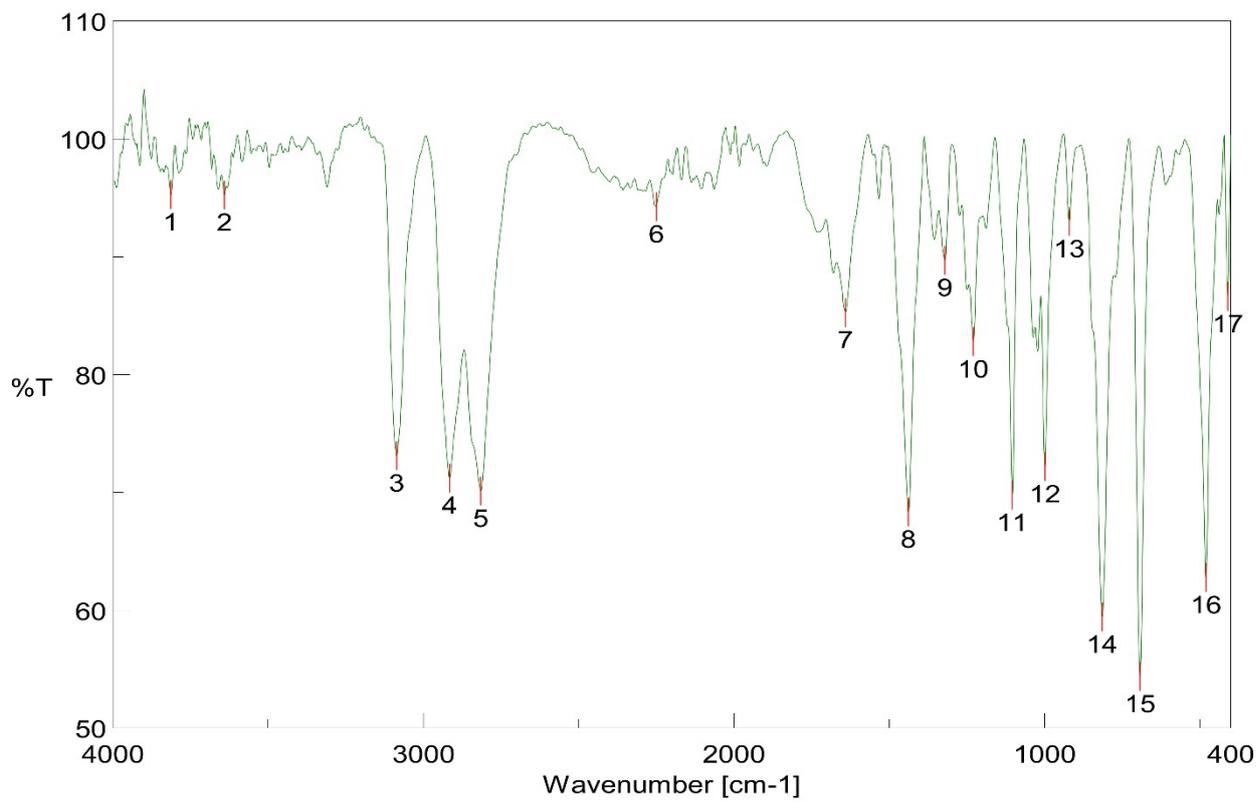


Fig. S14. FT-IR spectra of ligand L².

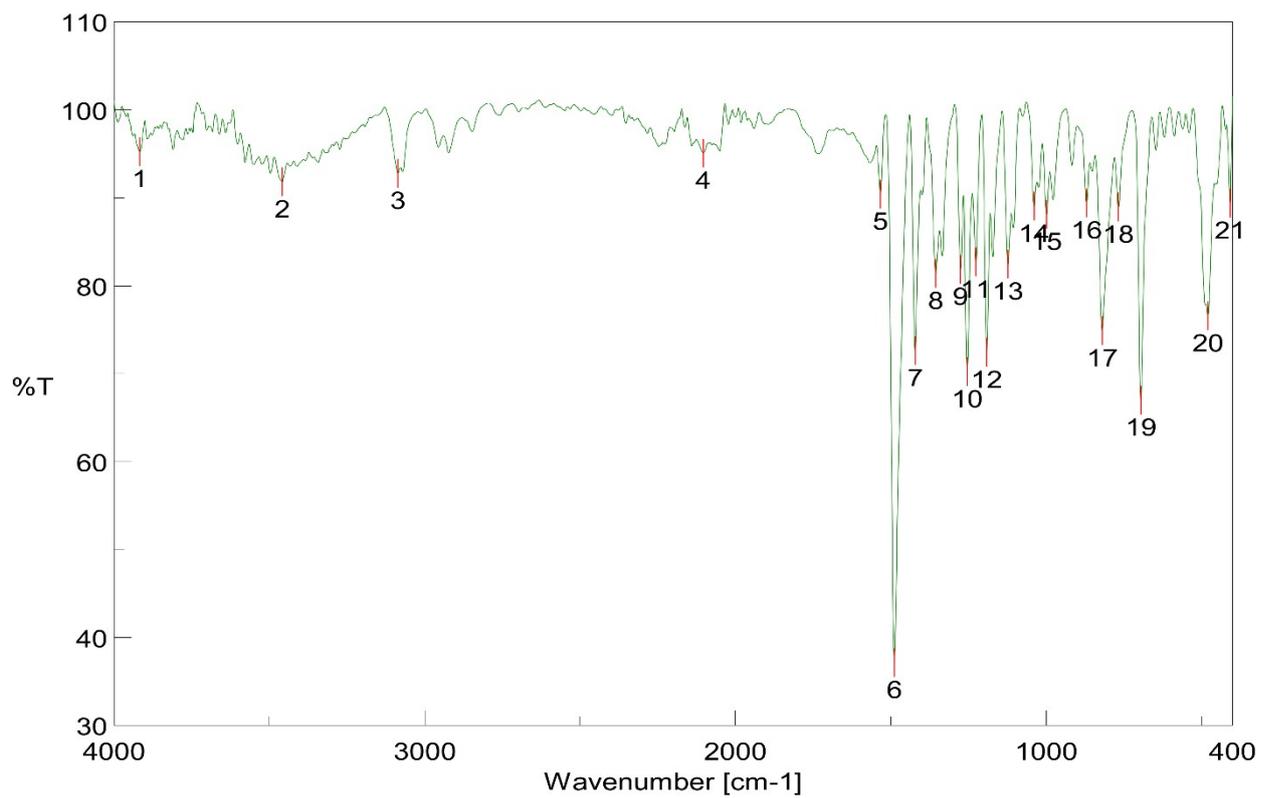


Fig. S15. FT-IR spectra of complex 1.

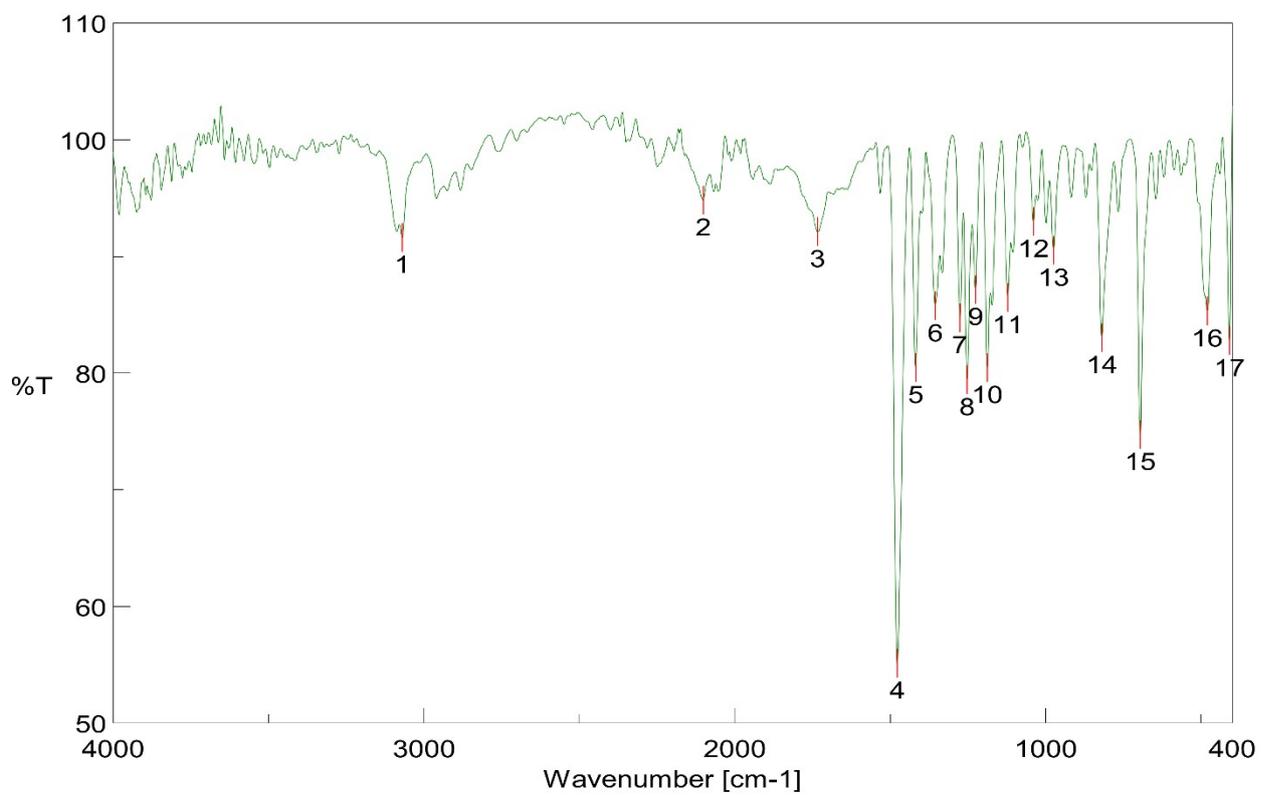


Fig. S16. FT-IR spectra of complex 2.

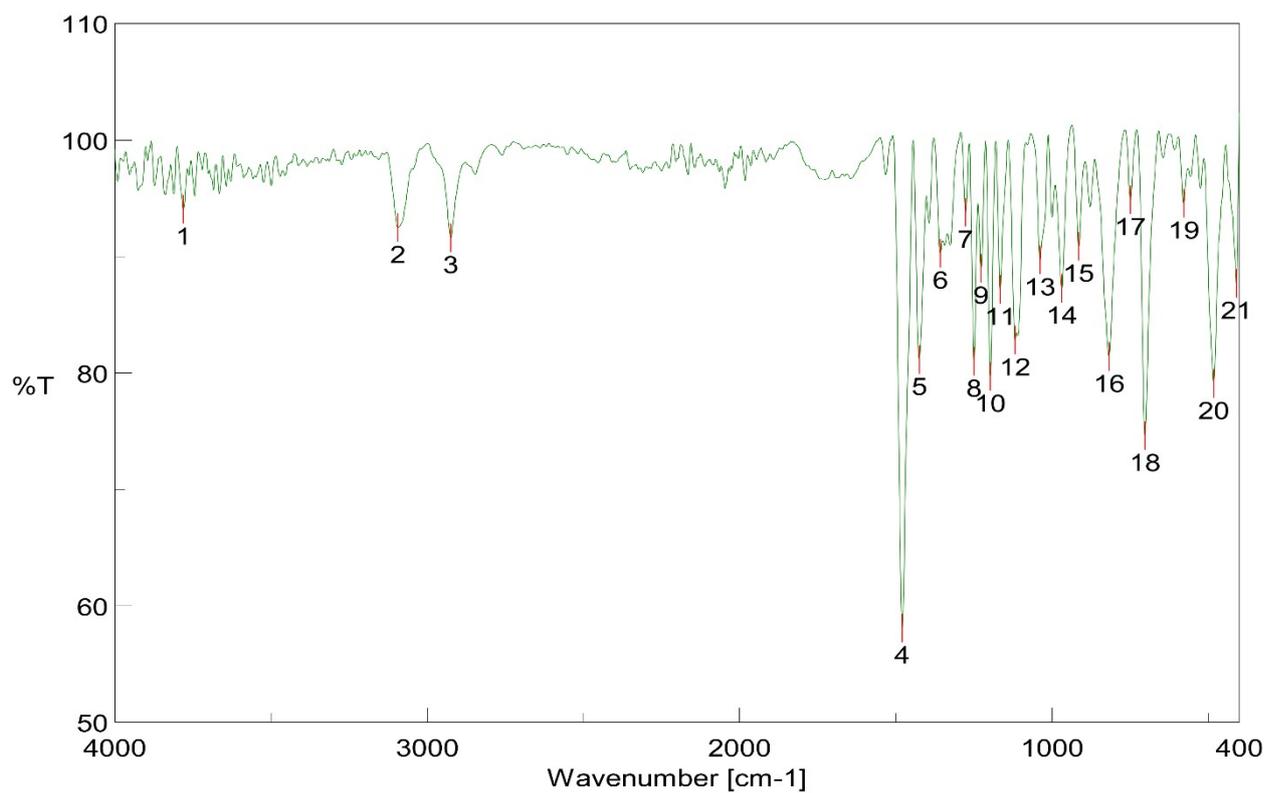


Fig. S17. FT-IR spectra of complex 3.

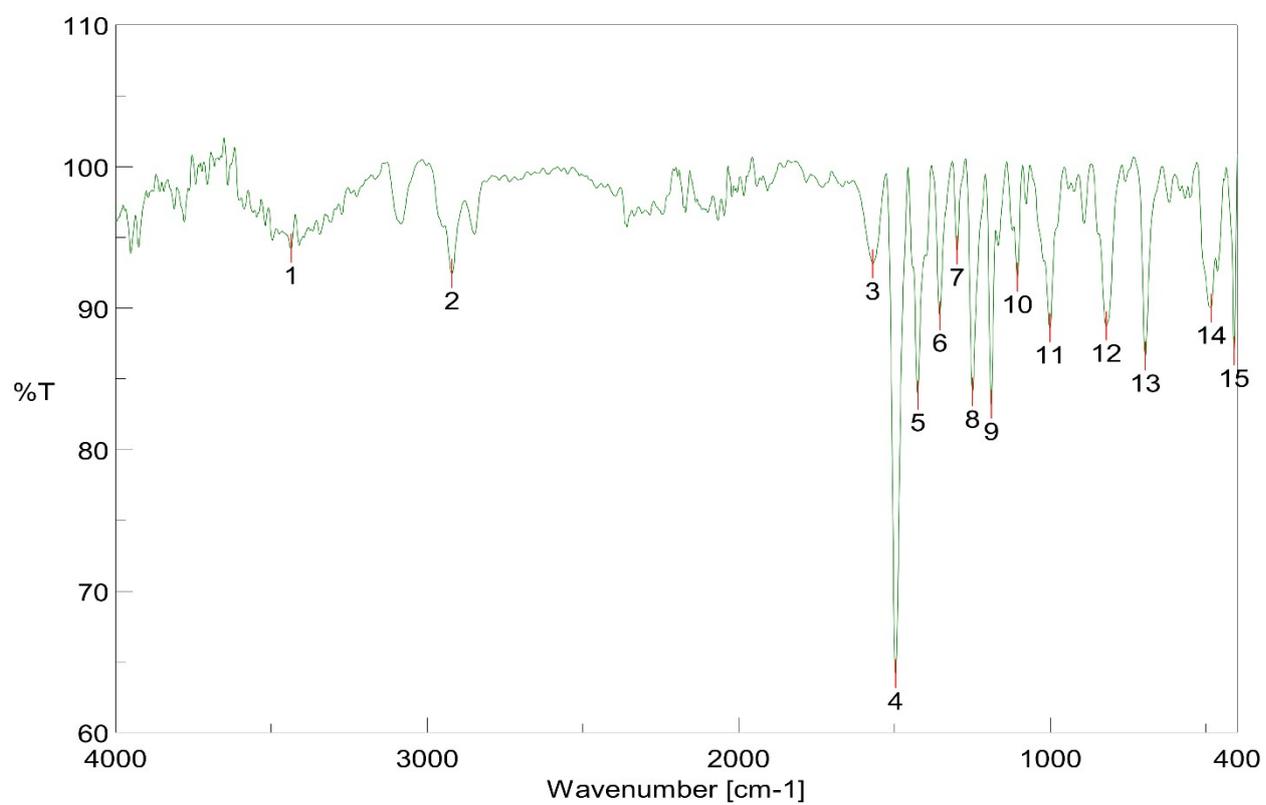


Fig. S18. FT-IR spectra of complex 4.

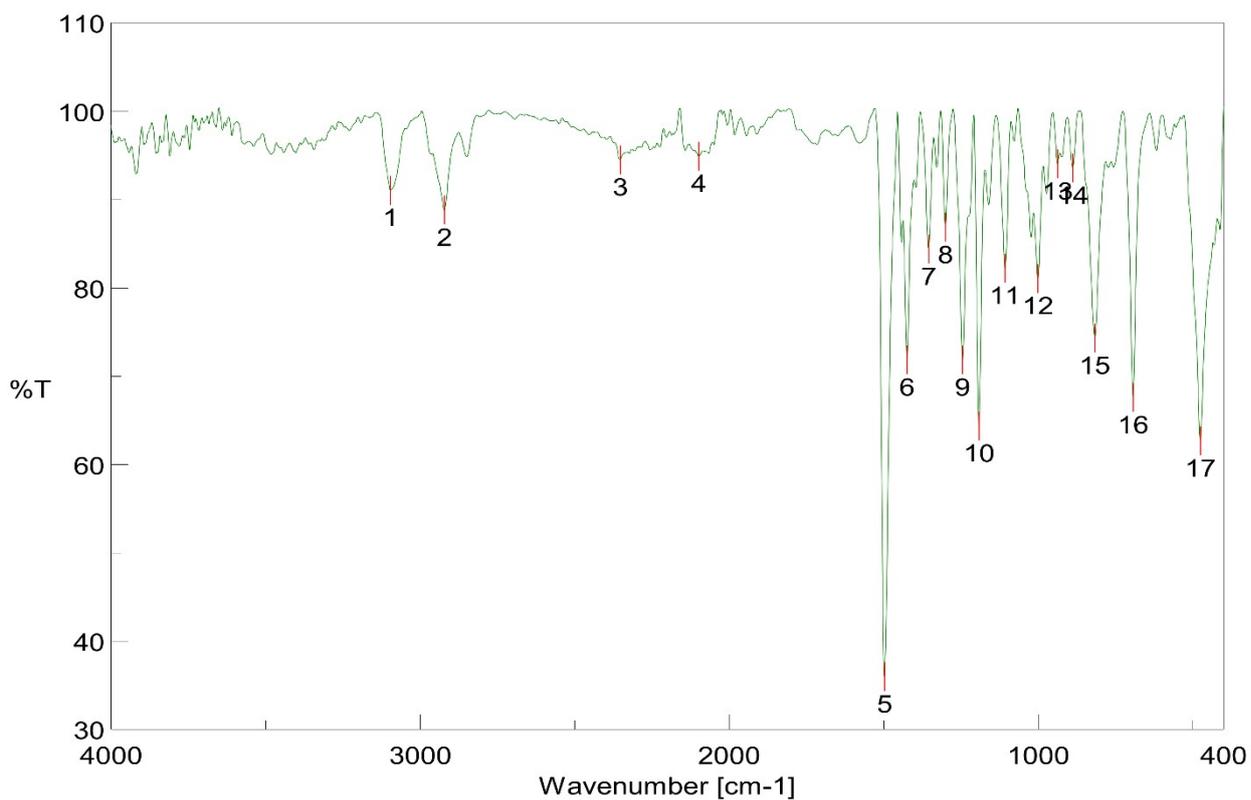


Fig. S19. FT-IR spectra of complex 5.

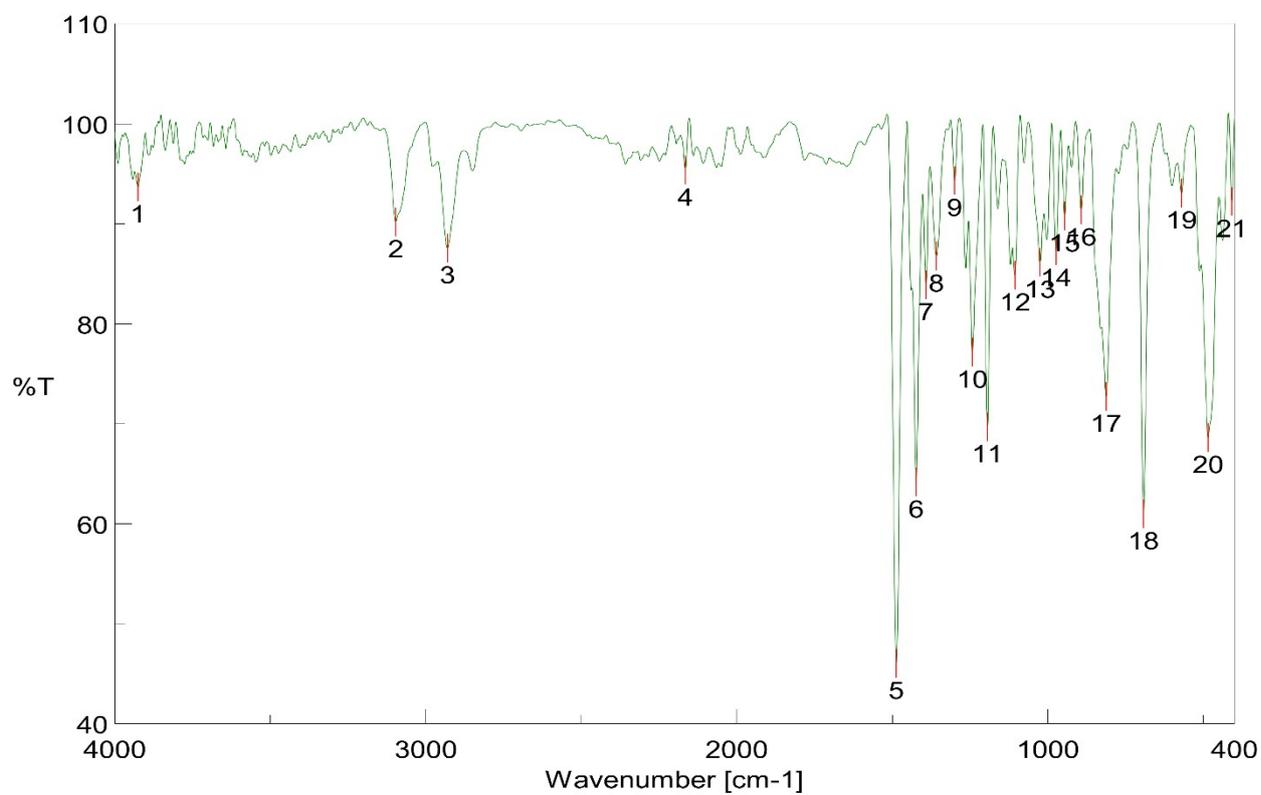


Fig. S20. FT-IR spectra of complex 6.

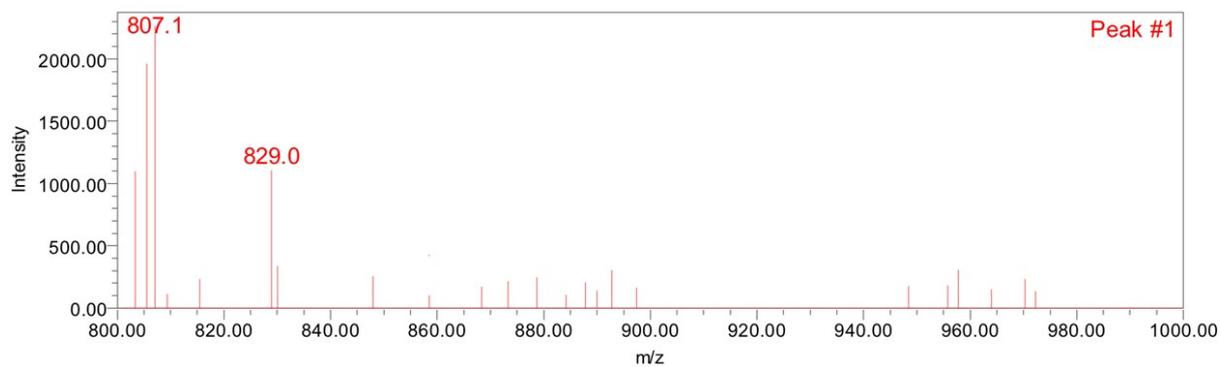


Fig. S21. ESI-MS spectra of complex 1.

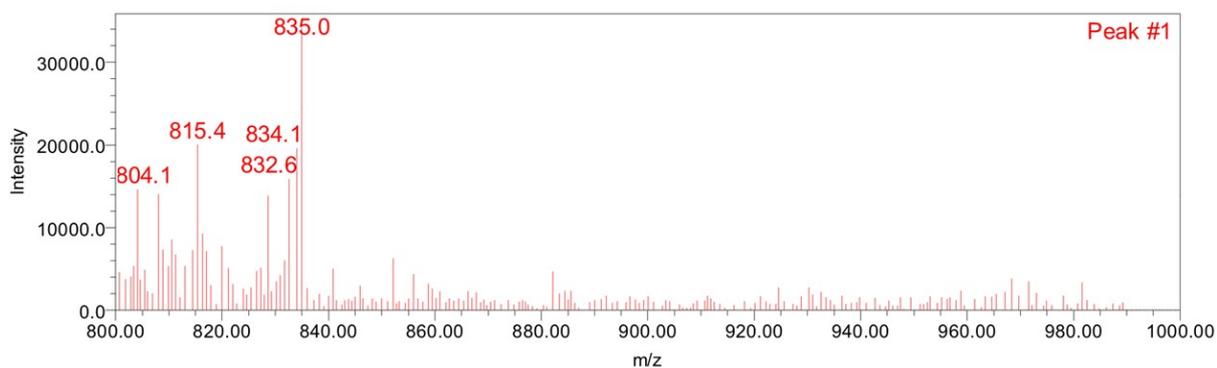


Fig. S22. ESI-MS spectra of complex 2.

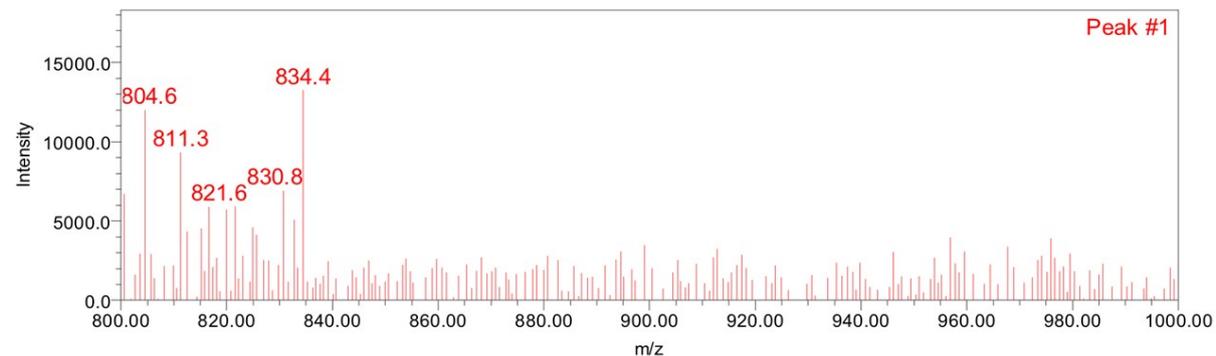


Fig. S23. ESI-MS spectra of complex 3.

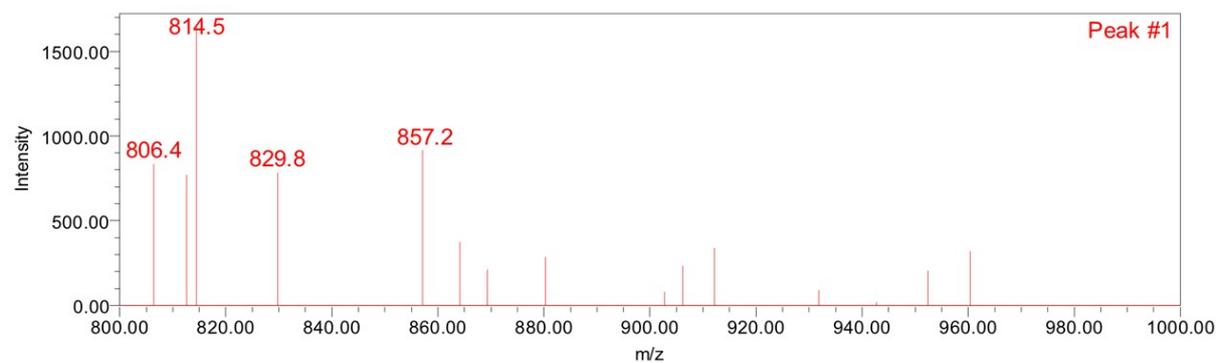


Fig. S24. ESI-MS spectra of complex 4.

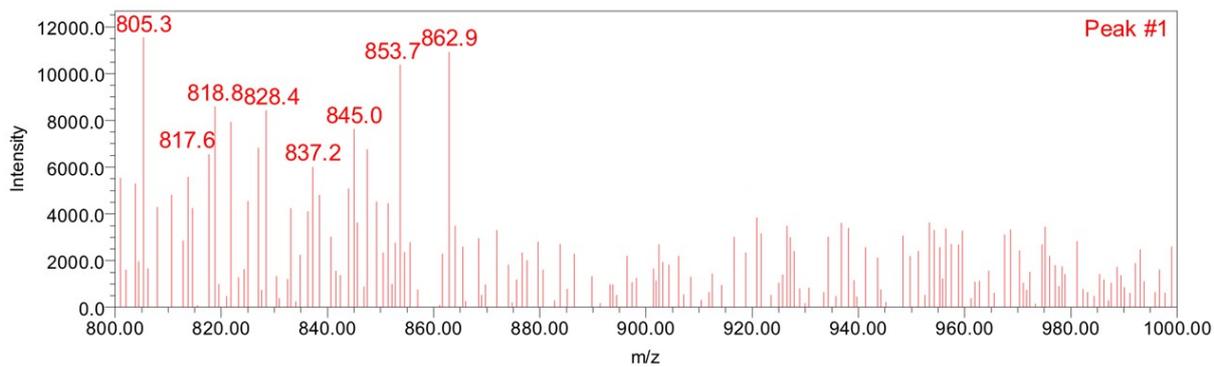


Fig. S25. ESI-MS spectra of complex 5.

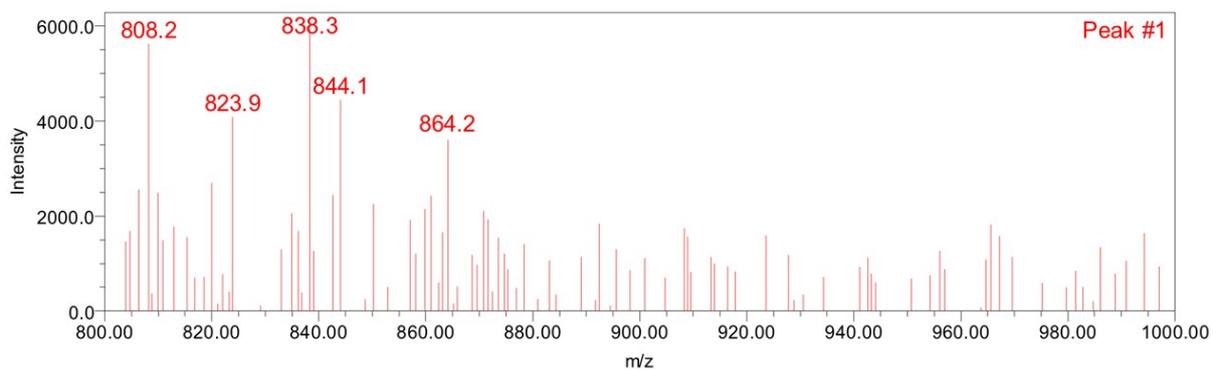


Fig. S26. ESI-MS spectra of complex 6.

Table S1. UV-visible spectral data of ligand precursors and complexes **1-6** in 10^{-5} DCM.

Entry	UV-vis spectral data (10^{-5} DCM) λ_{\max} nm (ϵ L mol $^{-1}$ cm $^{-1}$)	Wave Number (cm $^{-1}$)	Magnetic moment μ_{eff} (BM)
L¹	272 (08535) ($\pi \rightarrow \pi^*$)	36764	-
	306 (11321) ($n \rightarrow \pi^*$)	32679	
L²	270 (08338) ($\pi \rightarrow \pi^*$)	37037	-
	318 (03586) ($n \rightarrow \pi^*$)	31446	
1	278 (28414) ($\pi \rightarrow \pi^*$)	35971	dia
	332 (60925) ($n \rightarrow \pi^*$)	30120	
	394 (23557) charge transfer	25380	
	612 (01204) d-d	16339	
2	268 (63024) ($\pi \rightarrow \pi^*$)	37313	1.79
	334 (05759) ($n \rightarrow \pi^*$)	29940	
	436 (15283) charge transfer	22935	
	616 (02014) d-d	16233	
3	276 (68174) ($\pi \rightarrow \pi^*$)	36231	dia
	324 (9135) ($n \rightarrow \pi^*$)	30864	
4	270 (44459) ($\pi \rightarrow \pi^*$)	37037	dia
	332 (120532) ($n \rightarrow \pi^*$)	30120	
	398 (49608) charge transfer	25125	
	608 (0809) d-d	16447	
5	268 (114986) ($\pi \rightarrow \pi^*$)	37313	1.78
	308 (106755) ($n \rightarrow \pi^*$)	32467	
	434 (24588) charge transfer	23041	
	618 (02059) d-d	16181	
6	266 (195941) ($\pi \rightarrow \pi^*$)	37593	dia
	310 (86146) ($n \rightarrow \pi^*$)	32258	

Table S2. Summary of computational (DFT) data for complexes **1-6**.

Code	Energy (Hartree)	HOMO (eV)	LUMO (eV)	Band Gap (eV)	Theoretical λ_{\max} value (nm)
1	-4229.70	-5.650	-2.272	3.378	370
2	-4256.53	-5.592	-1.382	4.210	294
3	-4125.99	-5.642	-2.297	3.345	288
4	-4308.33	-5.732	-2.217	3.515	353
5	-4335.17	-5.524	-1.289	4.235	293
6	-4204.62	-5.825	-2.468	3.357	270

Table S3. Electrochemical parameters derived from the voltammograms for ligands **L¹**-**L²** and complexes (**1-6**).

Scan rate (mV/s)	$E_{p,a}$ (V)	$E_{p,c}$ (V)	$\Delta E = E_{p,a} - E_{p,c}$ (V)	$E^0 = (E_{p,a} + E_{p,c})/2$ (V)
L¹				
50	1.088	0.222	0.866	0.655
100	1.219	-0.051	1.270	0.584
150	1.352	-0.158	1.510	0.597
200	1.413	-0.234	1.647	0.589
1				
50	0.830	0.555	0.275	0.692
100	0.896	0.464	0.432	0.680
150	0.937	0.435	0.502	0.686
200	0.979	0.401	0.578	0.690
2				
50	1.040	0.316	0.724	0.678
100	1.287	-0.033	1.320	0.627
150	1.362	-0.099	1.461	0.631
200	1.369	-0.081	1.450	0.644
3				
50	1.030	0.377	0.653	0.703
100	1.213	0.214	0.999	0.713
150	1.245	0.089	1.156	0.667
200	1.194	-0.149	1.343	0.522
L²				
50	0.981	0.144	0.837	0.562
100	1.117	0.009	1.108	0.563
150	1.201	-0.071	1.272	0.565
200	1.272	-0.126	1.398	0.699
4				
50	0.871	0.578	0.293	0.724
100	0.624	0.361	0.263	0.492
150	0.604	0.348	0.256	0.476
200	0.557	0.329	0.228	0.443
5				
50	1.180	-0.016	1.196	0.582
100	1.342	-0.310	1.652	0.516
150	1.374	-0.387	1.761	0.493
200	1.288	-0.442	1.730	0.423
6				
50	0.746	0.426	0.320	0.586
100	0.774	0.394	0.380	0.584
150	0.858	0.366	0.492	0.612
200	0.911	0.330	0.581	0.620

Table S4. Thermogravimetric data of dithiocarbamate metal complexes **1-6**.

Entry	Steps	Temp.	Weight loss on TGA (%)	DTA peak (°C)	Inference
1	I	243-579	50.0	244.0	Mass loss continues even after 800 °C.
2	I	41-94	8.0	-	Mass loss continues even after 800 °C.
	II	95-428	54.0	183.0	
3	I	200-337	27.0	205.0	Mass loss continues even after 800 °C.
	II	338-480	6.0	-	
4	I	248-433	67.0	256.3	Mass loss continues even after 800 °C.
5	I	185-448	66.0	203.0	Mass loss continues even after 800 °C.
6	I	229-454	66.0	261.7	Mass loss continues even after 800 °C.

Table S5. Antiproliferative activity (GI_{50} , μM) against human cancer cell lines of synthesized ligands **L¹**, **L²**, and compounds **1-6**.

Compound	Antiproliferative activity (GI_{50}) ^a					
	Cell line (origin)					
	A549 (lung)	HeLa (cervix)	MIA PaCa- 2 (pancreas)	SW1573 (lung)	T-47D (breast)	WiDr (colon)
L¹	42 ± 13	22 ± 4.8	>100	88 ± 21	74 ± 29	52 ± 20
L²	83 ± 24	40 ± 7.5	77 ± 34	78 ± 33	30 ± 5.2	37 ± 6.5
1	>100	>100	>100	>100	>100	>100
2	>100	>100	>100	>100	>100	>100
3	12 ± 3.2	7.7 ± 0.74	7.5 ± 0.62	5.3 ± 1.1	66 ± 19	37 ± 13
4	>100	>100	>100	>100	>100	>100
5	>100	>100	>100	>100	>100	>100
6	11 ± 1.7	5.2 ± 0.62	7.5 ± 2.3	5.8 ± 0.07	>100	>100
CDDP ^c	4.9 ± 0.2	1.8 ± 0.5	n.d.	2.7 ± 0.4	17 ± 3.3	23 ± 4.3
5-FU ^c	2.2 ± 0.3	16 ± 4.5	n.d.	3.3 ± 1.2	43 ± 16	49 ± 6.7

Values represent mean ± standard deviation of at least three independent experiments. n.d. = not determined.

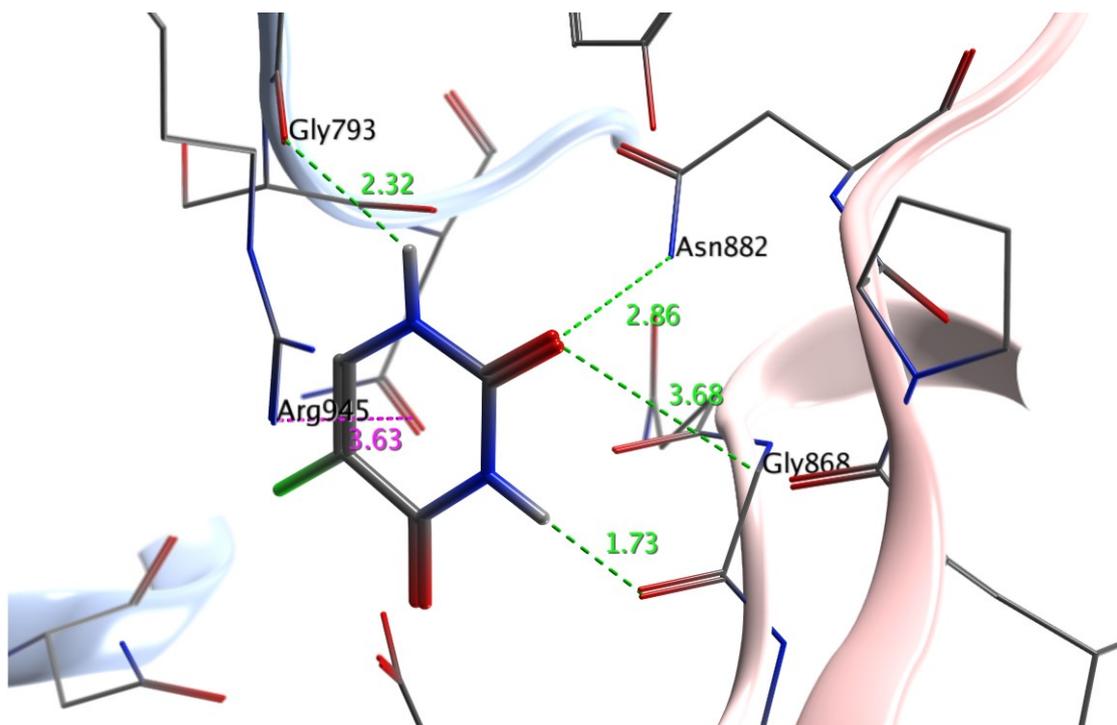


Fig. S27. Molecular docking studies of 5-fluorouracil (5-FU) with human topoisomerase II beta in complex with DNA (PDB ID: 3qx3).

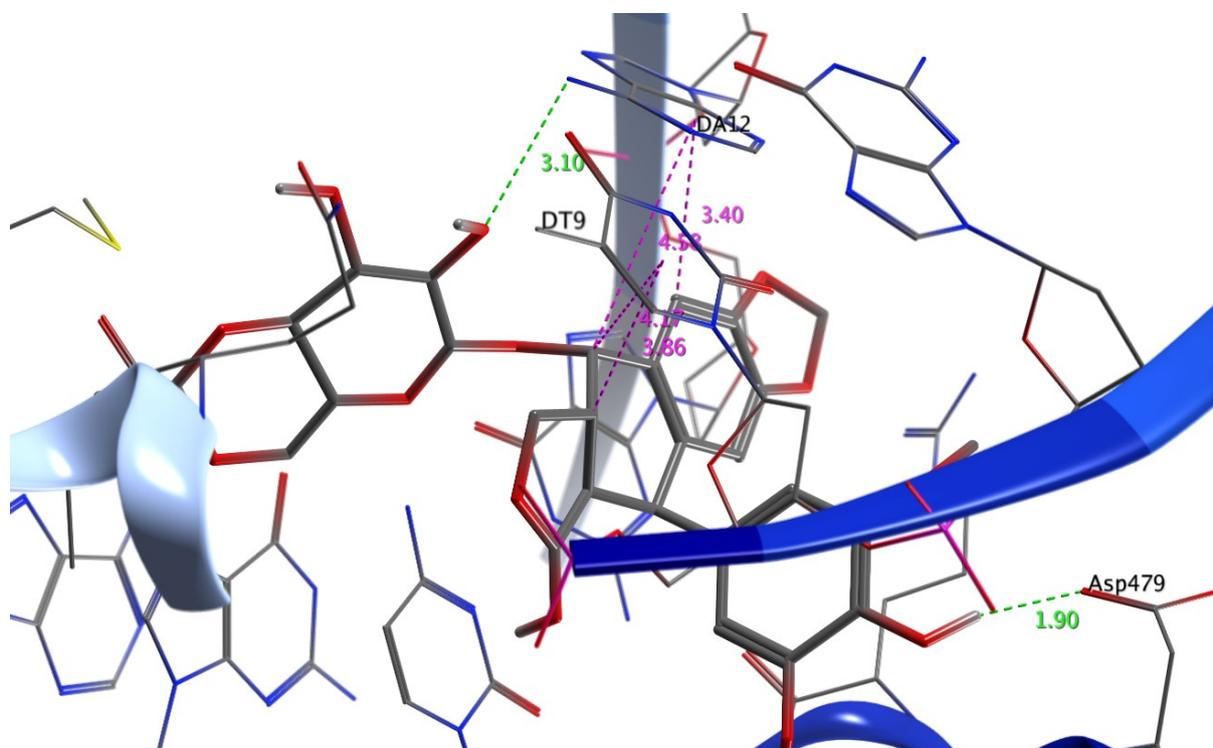


Fig. S28. Molecular docking studies of co-ligand (native) with human topoisomerase II beta in complex with DNA (PDB ID: 3qx3).

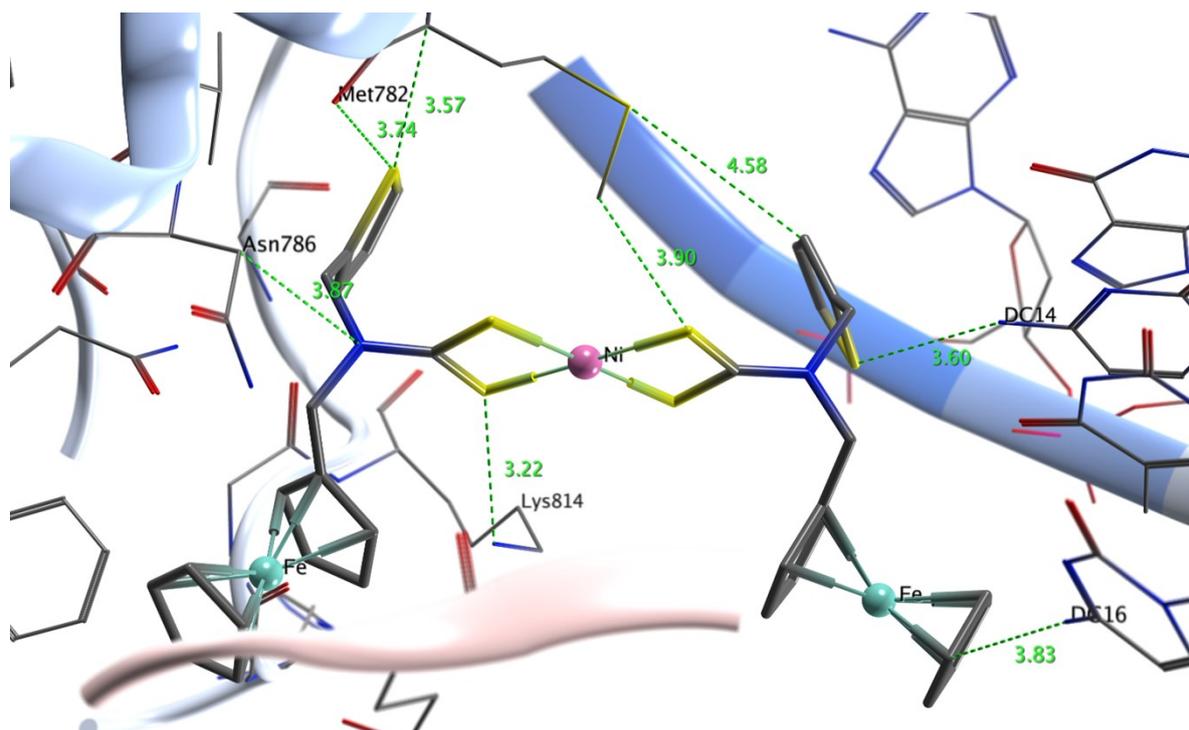


Fig. S29. Molecular docking studies of complex **1** with human topoisomerase II beta in complex with DNA (PDB ID: 3qx3).

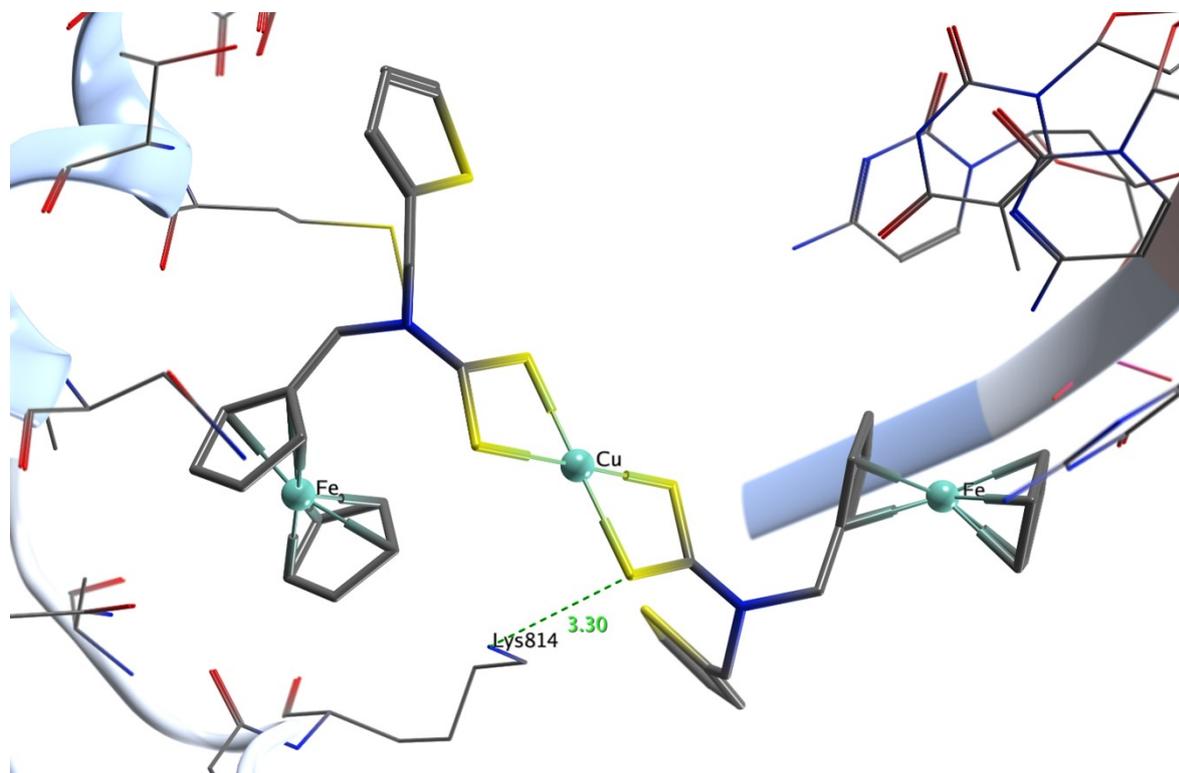
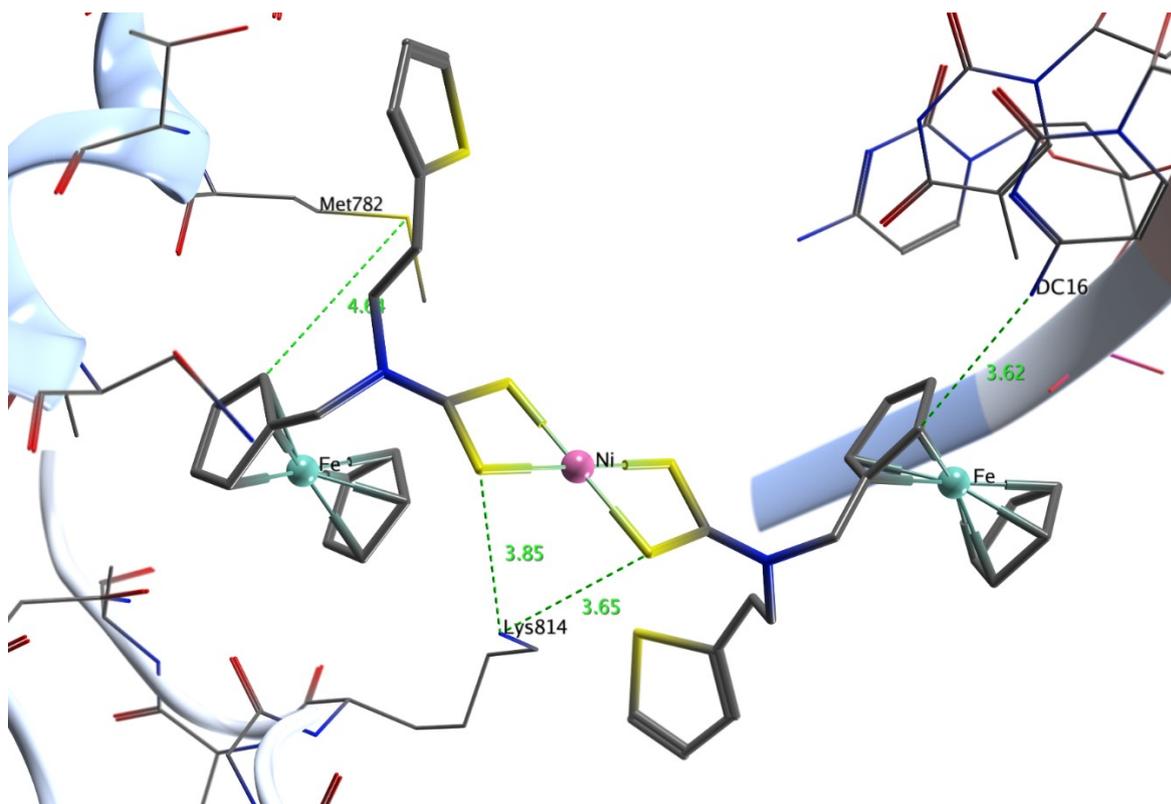


Fig. S30. Molecular docking studies of complex **2** with human topoisomerase II beta in complex with DNA (PDB ID: 3qx3).



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g. S31. Molecular docking studies of complex 4 with human topoisomerase II beta in complex with DNA (PDB ID: 3qx3).

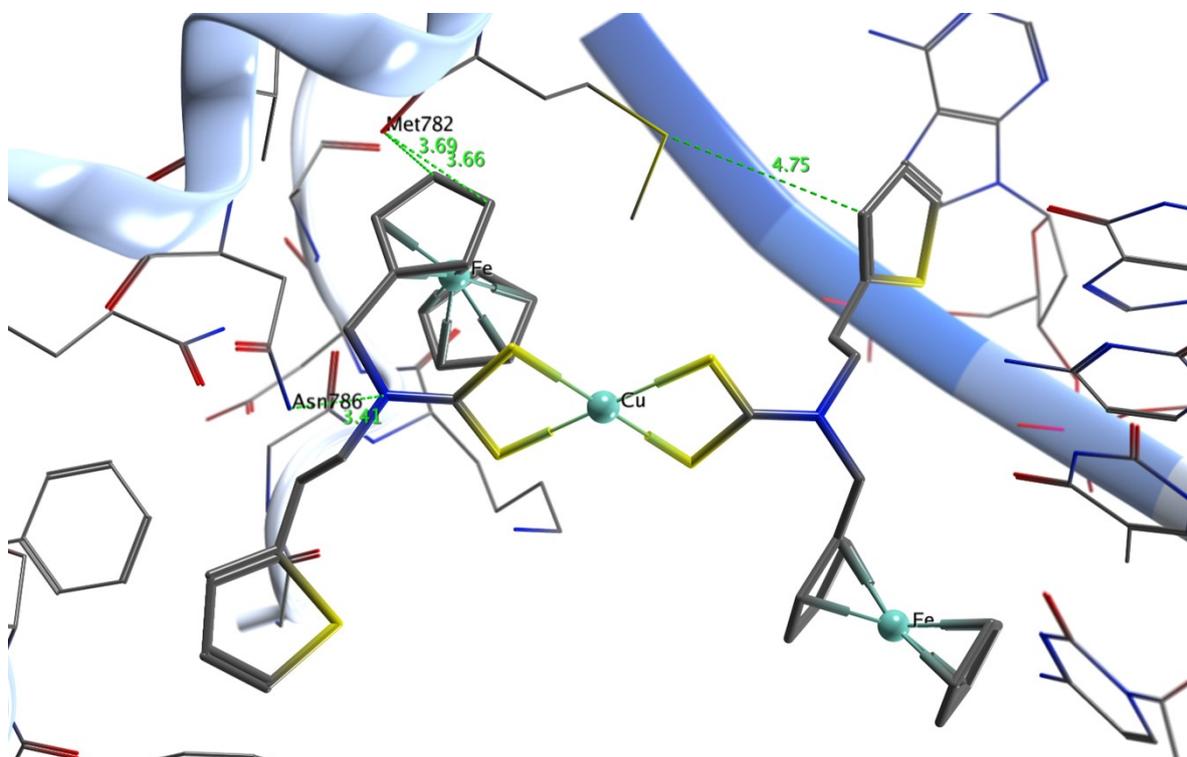


Fig. S32. Molecular docking studies of complex 5 with human topoisomerase II beta in complex with DNA (PDB ID: 3qx3).

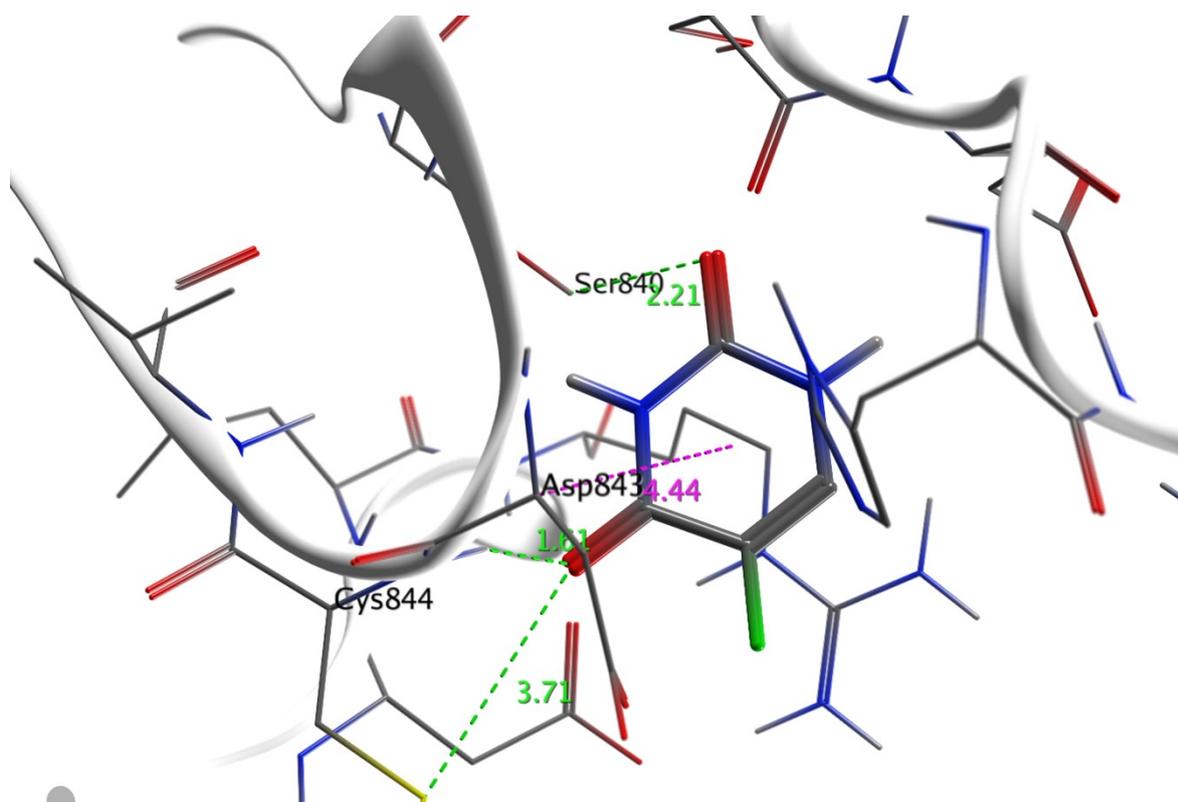


Fig. S33. Molecular docking studies of 5-fluorouracil (5-FU) with protein PI3K α (PDB ID: 4jps).

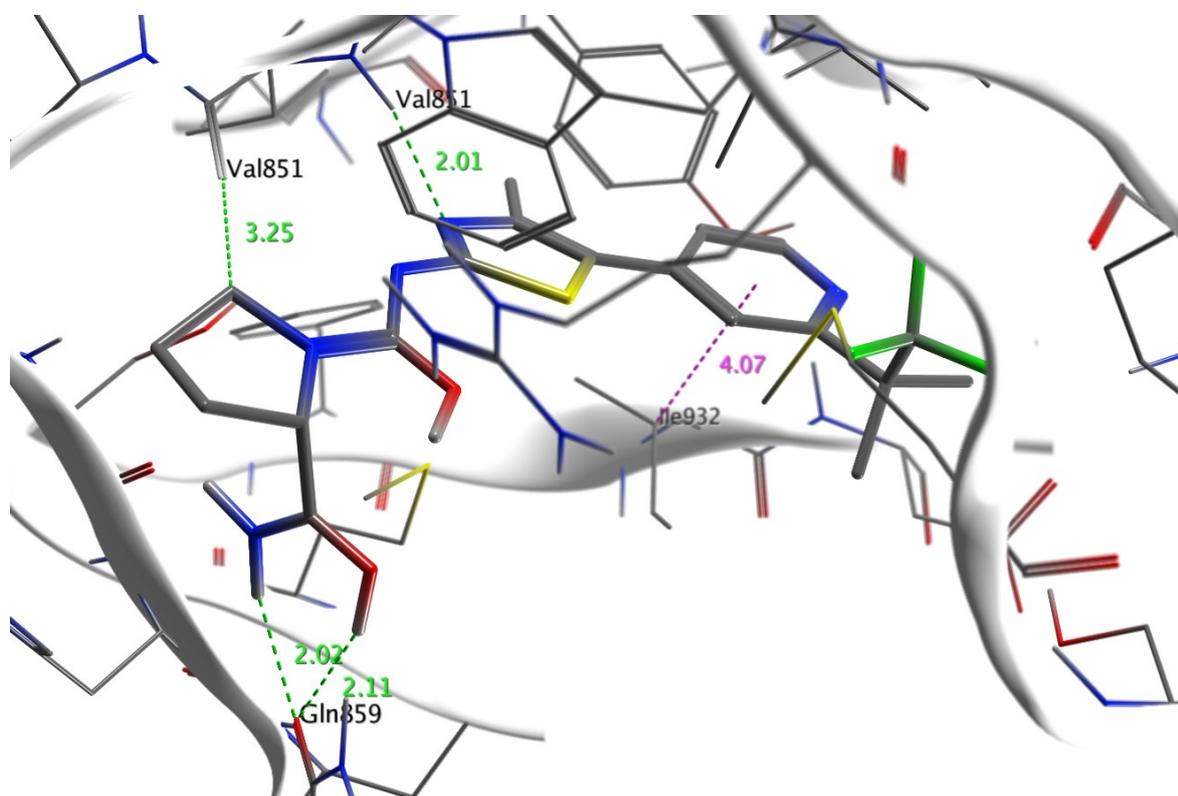


Fig. S34. Molecular docking studies of co-ligand (native) with protein PI3K α (PDB ID: 4jps).

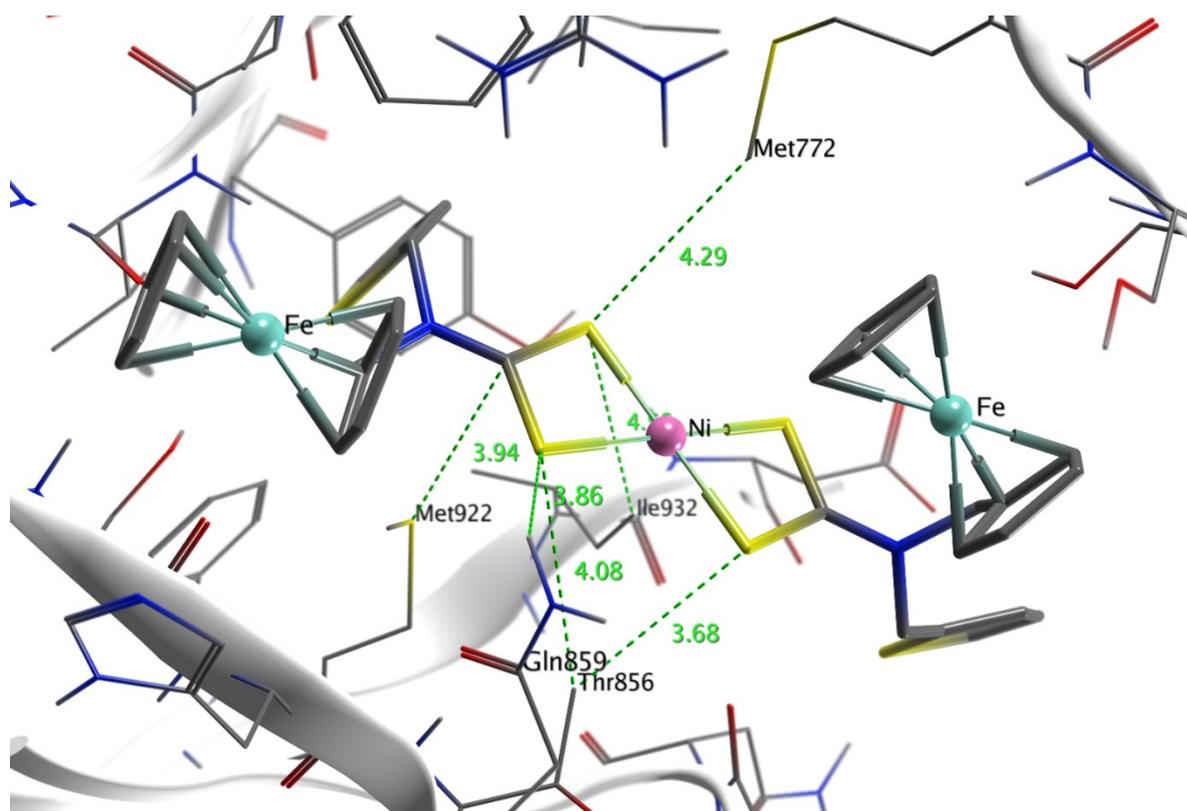


Fig. S35. Molecular docking studies of complex **1** with protein PI3K α (PDB ID: 4jps).

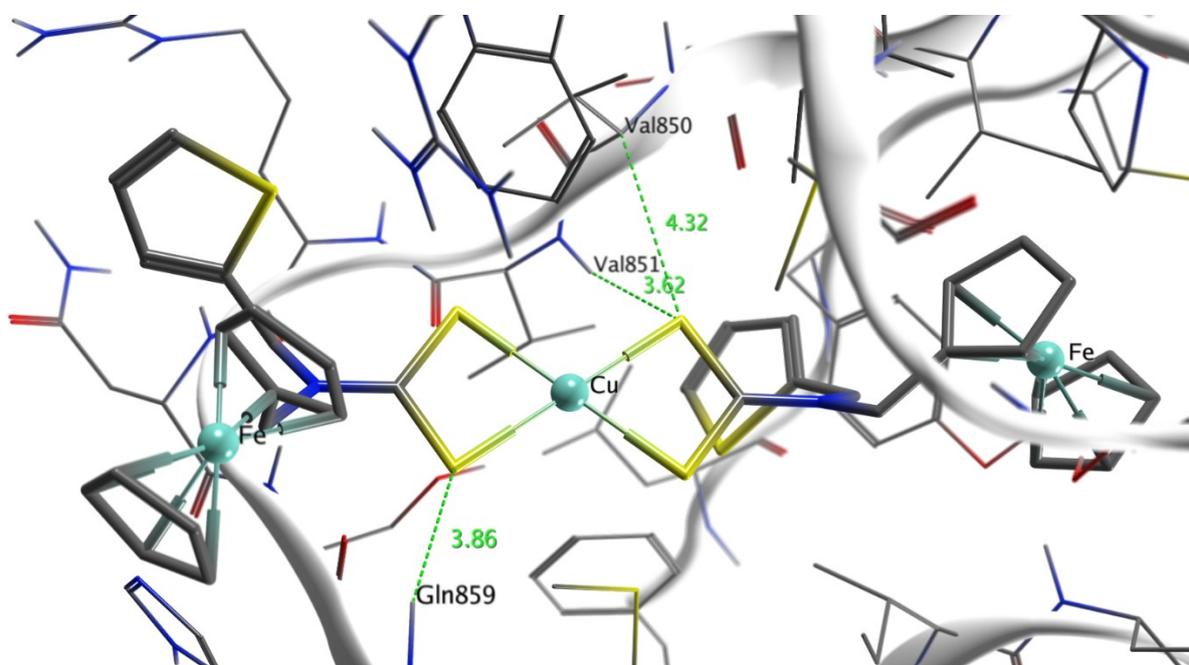


Fig. S36. Molecular docking studies of complex **2** with protein PI3K α (PDB ID: 4jps).

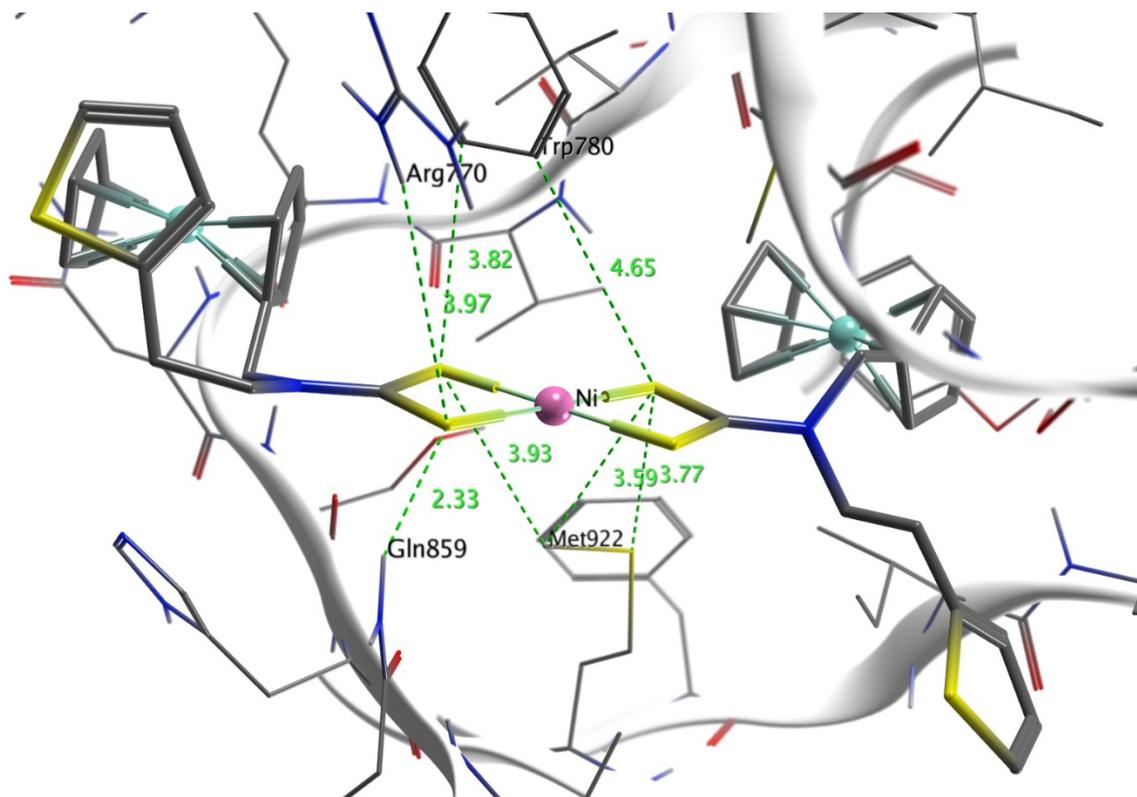


Fig. S37. Molecular docking studies of complex 4 with protein PI3K α (PDB ID: 4jps).

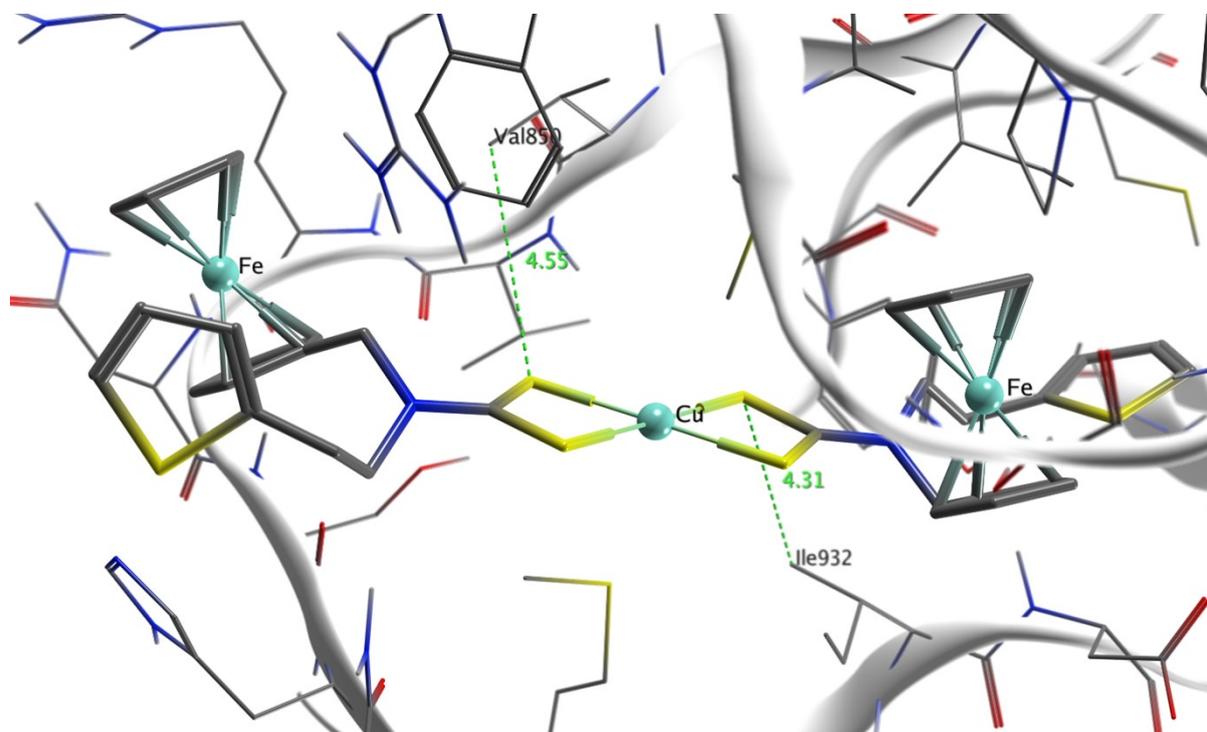


Fig. S38. Molecular docking studies of complex 5 with protein PI3K α (PDB ID: 4jps).