

Supporting Information

Synthesis, characterisation, molecular docking, biomolecular interaction and cytotoxicity studies of novel Ruthenium(II)-arene-2-heteroarylbenzoxazole complexes†

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Code	Absorption properties DMSO/DMSO-water		Emission properties DMSO/DMSO-water		Stokes shift (nm) DMSO/DMSO -water	Area DMSO/DMSO- water	Quantum yield DMSO/DMSO -water
	OD	(ϵ)	Excited (nm)	Emission (nm)			
5	0.46	6.5×10^3	307	--	--	1493	0.008
5W	0.71	1×10^5	312			365	0.001
5'	0.71	1×10^4	299	440	141	6272	0.02
5'W	0.38	5.4×10^3	338	413	75	1556	0.01
4	0.46	6.5×10^3	281	451	170	3300	0.02
4W	0.51	7.2×10^3	314	449	135	499	0.002
4'	0.46	6.5×10^3	313	440	127	8910	0.05
4'W	0.41	5.8×10^3	312	441	129	541	0.003
4'	0.45	6.4×10^3	270	441	171	7062	0.04
Ref	0.4	5.7×10^3	496	515	19	152890	0.95

Table S1. Photophysical properties of **4**, **4'**, **5** and **5'** benzoxazole based Ru(II) complexes

*W indicate DMSO: water (1:1)

Table S2. Physical characteristics of complexes

Complex	Chemical formula	Mol. wt	Solubility [mg/ml]
4	$C_{22}H_{21}BrClF_6N_2OPRu^+$	690.81	7
5	$C_{18}H_{13}BrClF_6N_2OPRu^+$	634.70	6
4'	$C_{26}H_{23}BrClF_6N_2OPRu^+$	740.87	8
5'	$C_{22}H_{15}BrClF_6N_2OPRu^+$	684.76	7

Table S3. Molar conductivity study of the complexes

Complex	Concentration [M]	Molar conductivity (Λ_m) ($\mu S/cm$)			
		DMF	10% aqueous DMF	DMSO	10% aqueous DMSO
4	5×10^{-5}	40	45	44	55
5	5×10^{-5}	41	45	41	50
4'	5×10^{-5}	19	63	22	32
5'	5×10^{-5}	65	81	36	76



Table S4. Binding energy of DNA-complexes and BSA-complexes calculated from Molecular docking studies (kcal mol⁻¹)

Complex	DNA Binding free energy ($\Delta G_{\text{binding}}$) ^a	BSA Binding free energy ($\Delta G_{\text{binding}}$) ^a
4	-8.2	-7.9
5	-8.7	-8.3
4'	-8.3	-7.7
5'	-8.9	-8.1

Table S5. Relative viscosity of CT-DNA interaction with Ru(II) based complex (**4**)

[Complex]/[DNA]	t1	t2	t3	Average(t)	(t-t0)	(η/η_0) ^{1/3}
Buffer only	132.02	131.47	132.21	131.9		
Buffer+DNA	138.61	138.56	138.76	138.64	6.74	1
2 μ M	139.5	139.69	140.03	139.74	7.84	1.05
4 μ M	140.65	140.92	140.87	140.81	8.91	1.09
6 μ M	141.23	141.64	141.79	141.55	9.65	1.12
8 μ M	142.68	142.98	142.7	142.78	10.88	1.17
10 μ M	144.01	144.23	144.17	144.14	12.23	1.22

Table S6. Relative viscosity of CT-DNA interaction with Ru(II) based complex (**5**)

[Complex]/[DNA]	t1	t2	t3	Average(t)	(t-t0)	(η/η_0) ^{1/3}
Buffer only	132.02	131.47	132.21	131.9		
Buffer+DNA	138.61	138.56	138.76	138.64	6.74	1
2 μ M	138.92	139.01	138.86	138.93	7.03	1.01
4 μ M	139.03	139.24	139.39	139.22	7.32	1.03
6 μ M	139.75	139.97	140.07	139.93	8.03	1.06
8 μ M	140.39	140.54	140.83	140.58	8.68	1.08
10 μ M	141.19	141.49	142	141.56	9.66	1.13

Table S7. Relative viscosity of CT-DNA interaction with Ru(II) based complex (**4'**)

[Complex]/[DNA]	t1	t2	t3	Average(t)	(t-t0)	(η/η_0) ^{1/3}
Buffer only	132.02	131.47	132.21	131.9		
Buffer+DNA	138.61	138.56	138.76	138.64	6.74	1
2 μ M	142.18	142.59	142.92	142.56	10.66	1.16
4 μ M	143.33	143.43	143.67	143.47	11.57	1.19
6 μ M	144.76	144.98	145.01	144.91	13.01	1.25
8 μ M	145.78	145.72	145.91	145.80	13.90	1.27
10 μ M	146.18	146.47	146.95	146.53	14.63	1.29

Table S8. Relative viscosity of CT-DNA interaction with Ru(II) based complex (**5'**)

[Complex]/[DNA]	t1	t2	t3	Average(t)	(t-t0)	(η/η_0) ^{1/3}
Buffer only	132.02	131.47	132.21	131.9		
Buffer+DNA	138.61	138.56	138.76	138.64	6.74	1
2 μ M	143.24	143.33	143.38	143.31	11.41	1.19
4 μ M	145.01	145.23	145.32	145.18	13.28	1.25
6 μ M	146.27	146.39	146.65	146.43	14.53	1.29
8 μ M	147.31	147.33	147.49	147.37	15.47	1.32
10 μ M	149.01	149.43	149.19	149.21	17.31	1.37

Table S9. Relative viscosity of CT-DNA interaction with Ru(II) based complex (**EtBr**)

[EtBr]/[DNA]	t1	t2	t3	Average(t)	(t-t0)	(η/η_0) ^{1/3}
Buffer only	132.02	131.47	132.21	131.9		
Buffer+DNA	138.61	138.56	138.76	138.64	6.74	1
2 μ M	144.85	145.06	144.92	144.943	13.04	1.25
4 μ M	146.19	146.23	146.31	146.243	14.34	1.29
6 μ M	147.12	147.38	147.92	147.473	15.57	1.32
8 μ M	148.03	148.62	148.95	148.533	16.63	1.35
10 μ M	149.89	150.14	150.17	150.067	18.2	1.4

Table S10. Selected bond lengths (Å) and bond angles (°) of complexes (**4**, **4'**, **5** and **5'**).

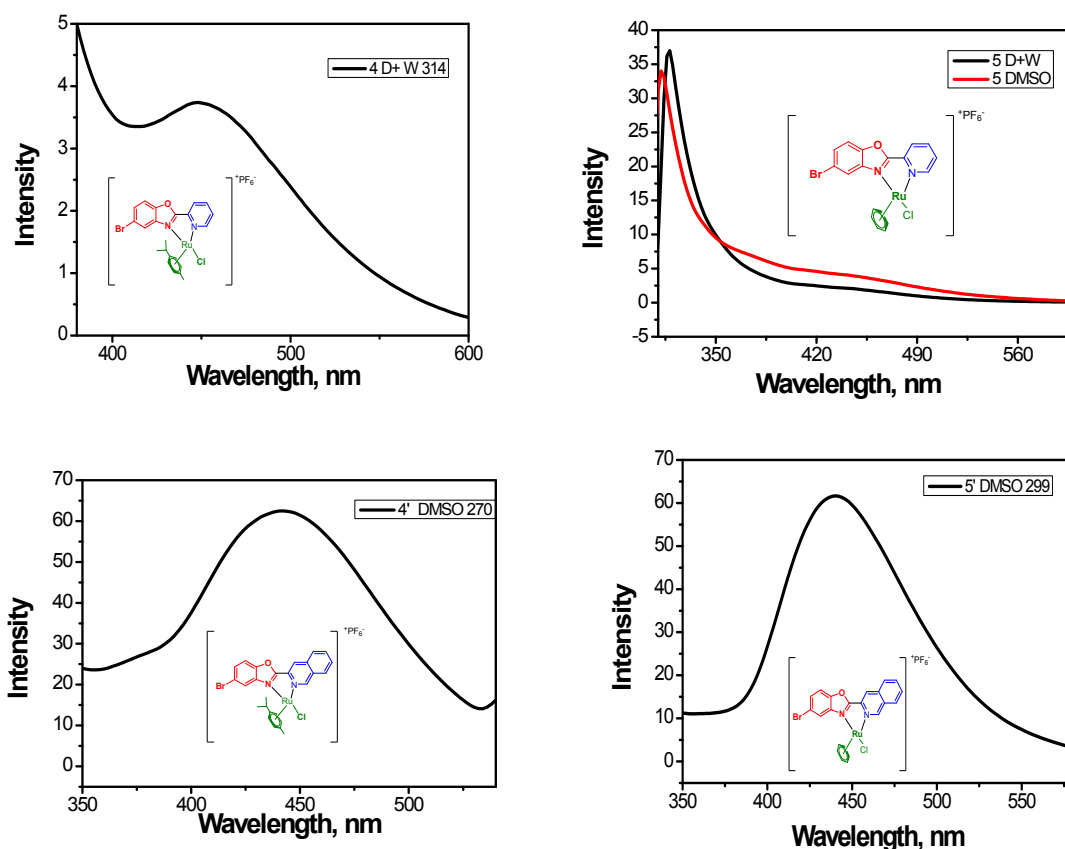
	4	5	4'	5'
Ru(1)–N(1)	2.0944	2.0944	2.0944	2.0944
Ru(1)–N(2)	2.1146	2.1146	2.1146	2.1146
Ru(1)–Cl(1)	2.399	2.3999	2.3999	2.3999
Ru(1)–Centroid*	2.2660	2.3678	2.1549	2.2328
N(1)–Ru(1)–N(2)	76.673	76.943	77.994	77.618
N(1)–Ru(1)–Cl(1)/Br(1)	82.593	84.711	87.334	87.083
N(2)–Ru(1)–Cl(1)/Br(1)	81.536	82.937	86.967	85.122

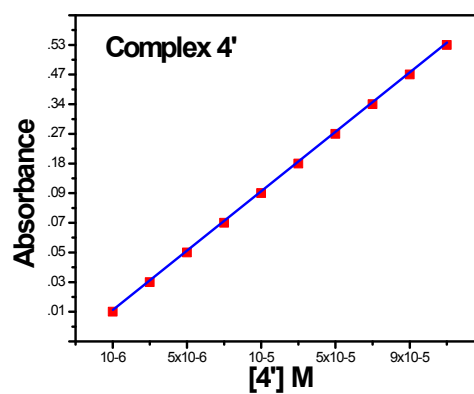
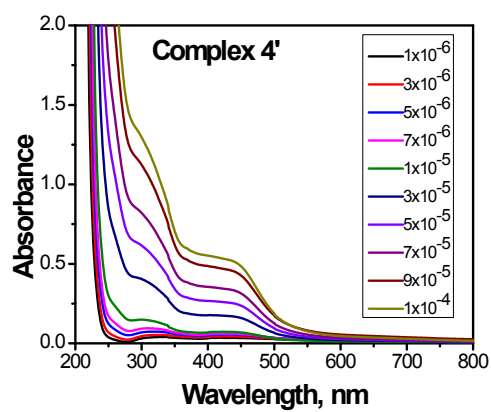
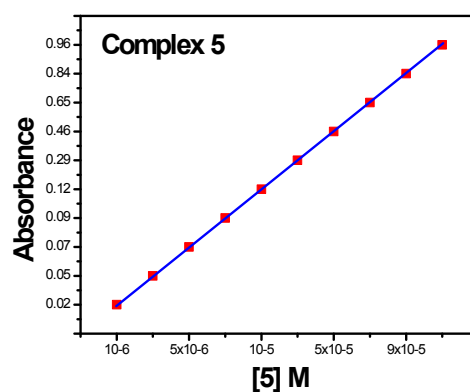
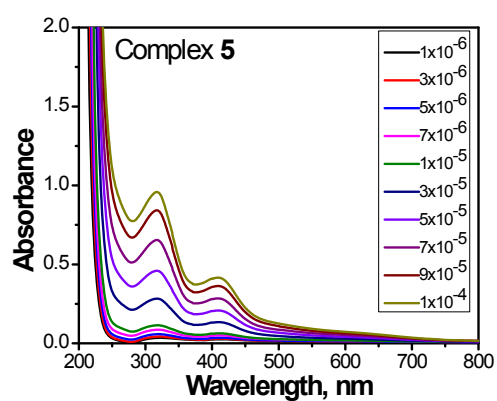
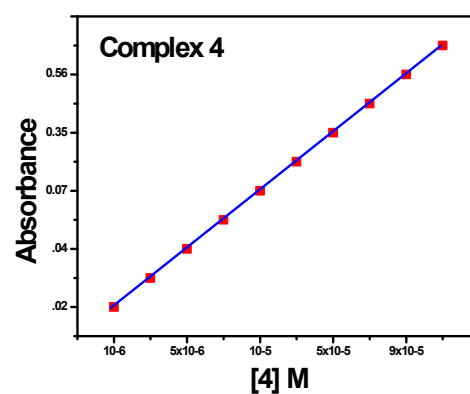
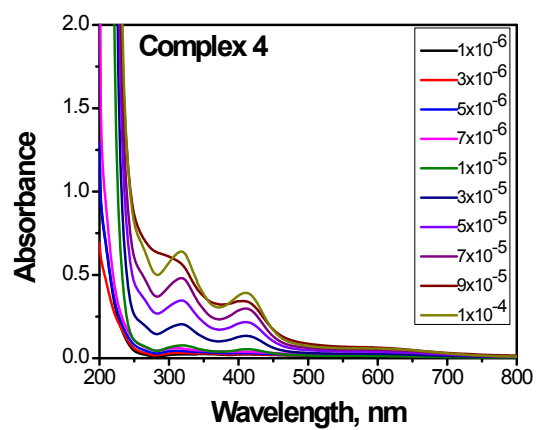
* N(1)= Benzoxazole nitrogen , N(2)= Pyridine nitrogen or quinioline nitrogen

Table S11. Calculated molecular electronic parameters.

Complex	Energy (kJ/mol)	DM (Debye)	E _H (eV)	E _L (eV)	ΔE (eV)	χ (eV)	η (eV)	μ (eV)	ω (eV)	S (eV)
4	-2514237.39	9.10	-8.89	-5.84	3.05	7.36	-1.53	-7.37	-17.78	-0.763
5	-2612936.52	8.33	-9.50	-6.52	2.98	8.01	-1.50	-8.10	-21.47	-0.747
4'	-2709349.02	7.75	-9.45	-6.84	2.61	8.14	-1.31	-8.14	-25.40	-0.653
5'	-2610624.73	9.15	-9.35	-6.71	2.64	8.03	-1.32	-8.03	-24.45	-0.660

Dipole moment (DM), Energy of HOMO (E_H), energy of LUMO (E_L), energy band gap (ΔE), electronegativity (χ), global hardness (η), chemical potential (μ), global electrophilicity index (ω) and global softness (S).

**Fig. S1** Emission spectra of **4**, **5**, **4'** and **5'** in DMSO and DMSO: water media



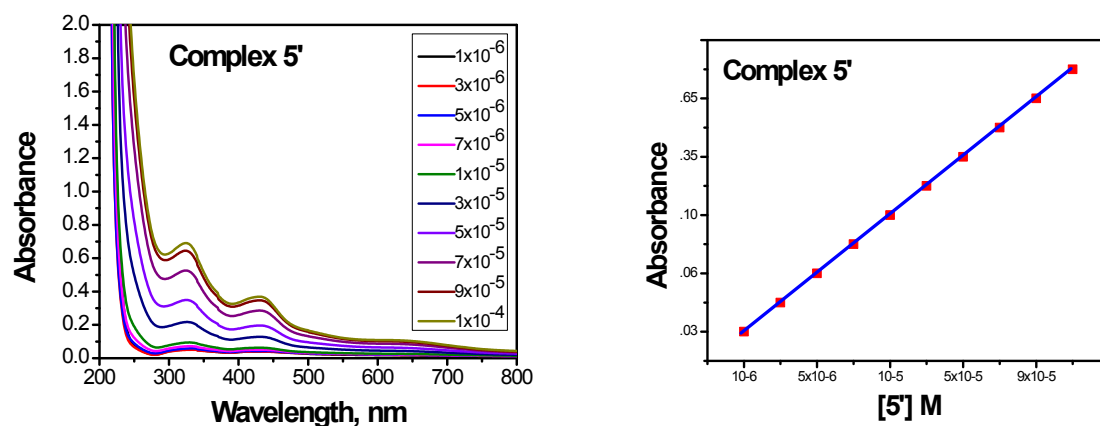
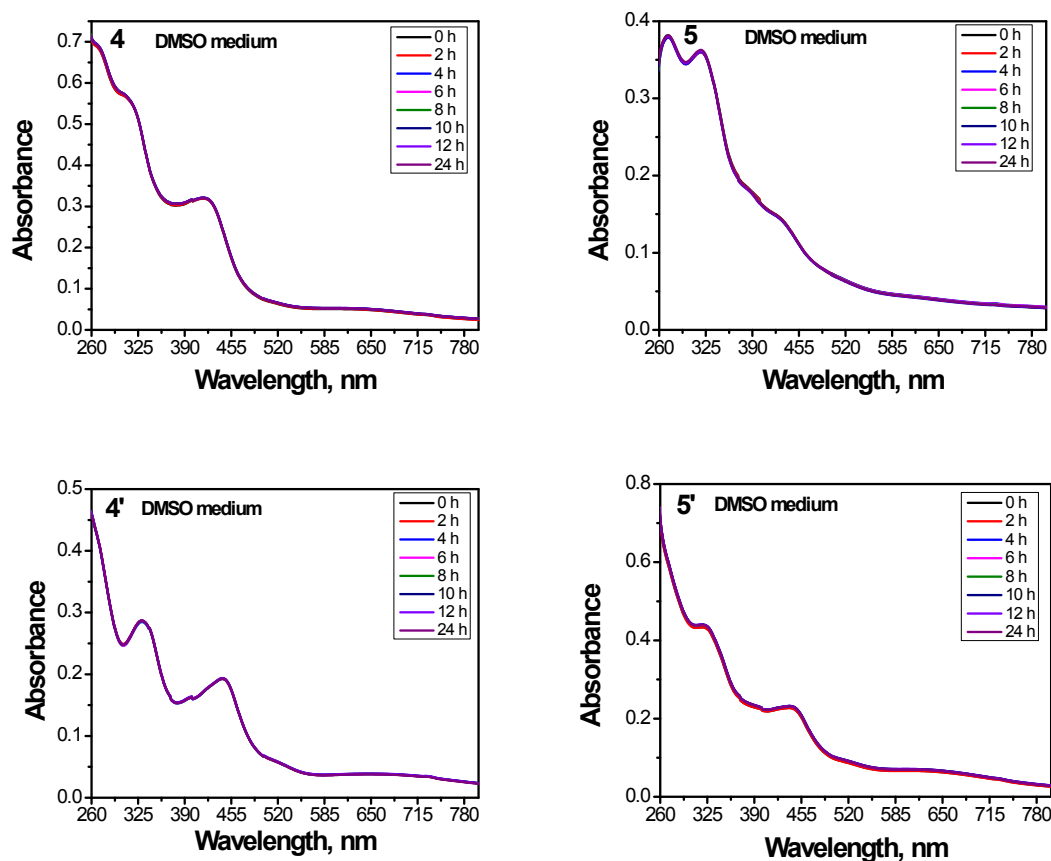
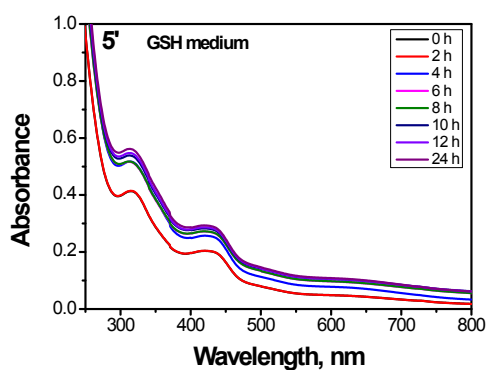
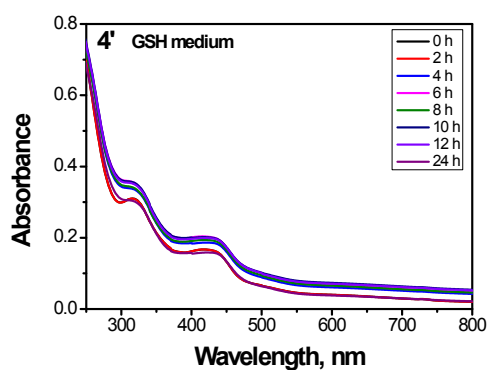
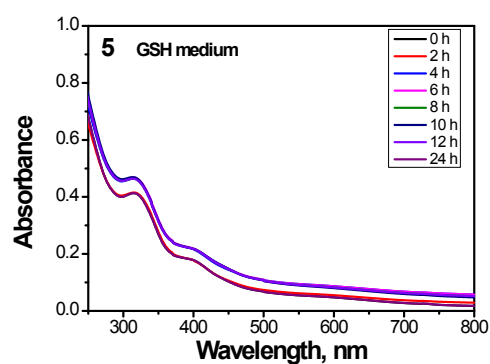
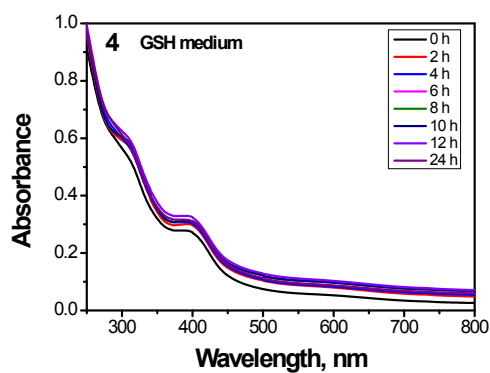


Fig. S2 Solubility study of 4, 5, 4' and 5' in DMSO: water media

(a) Stability in DMSO medium



(b) Stability in GSH medium



(c) Stability in Water medium

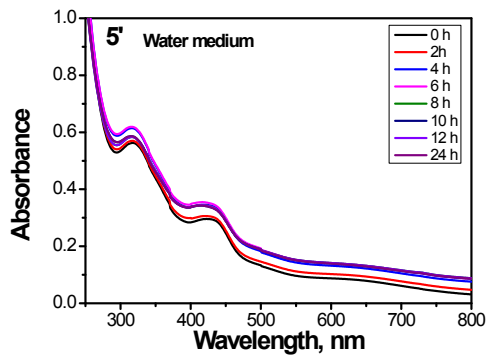
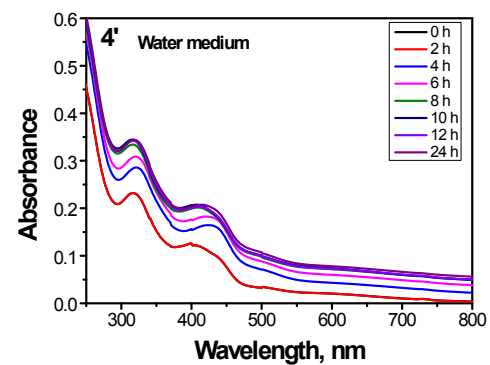
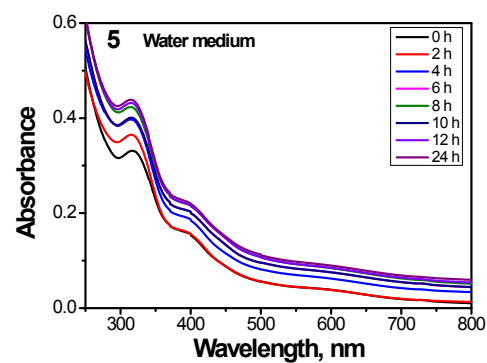
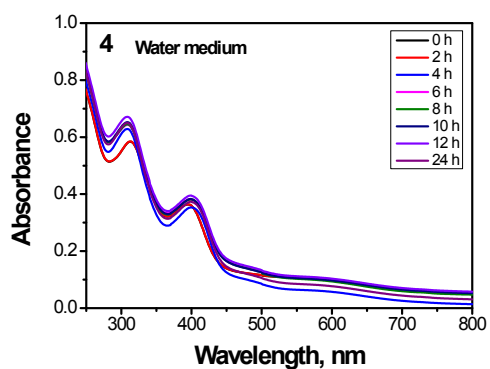


Fig. S3 Stability study of compound **4**, **5**, **4'** and **5'** complex in (a) DMSO (b) GSH (c) Water

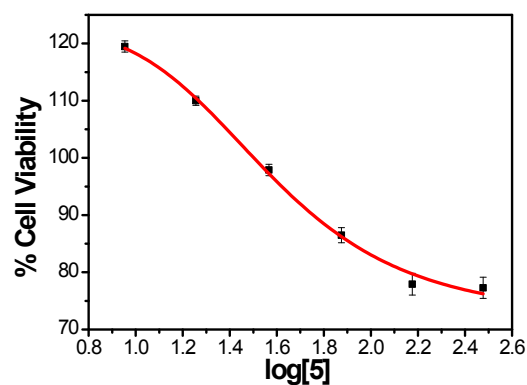
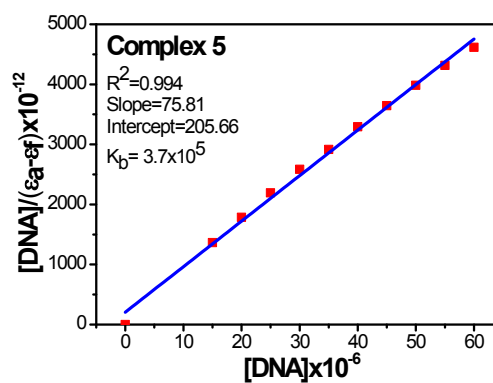
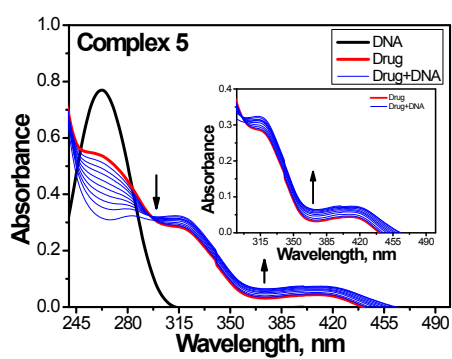
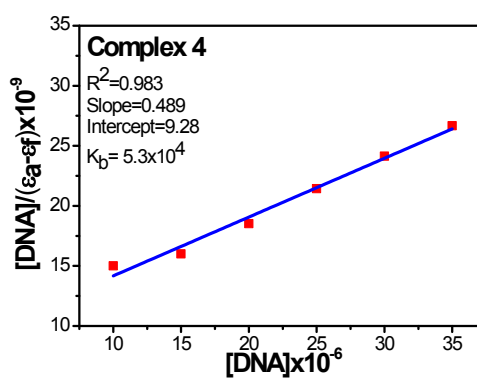
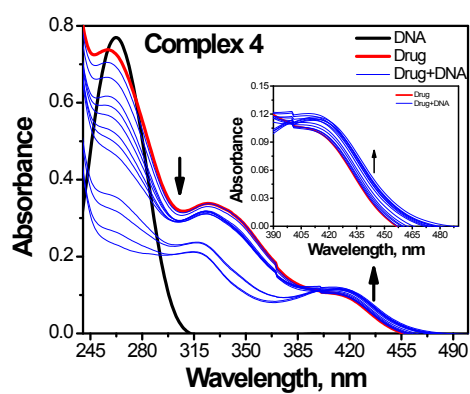


Fig. S4 Cell viability plot of complex **5** in HeLa cell line



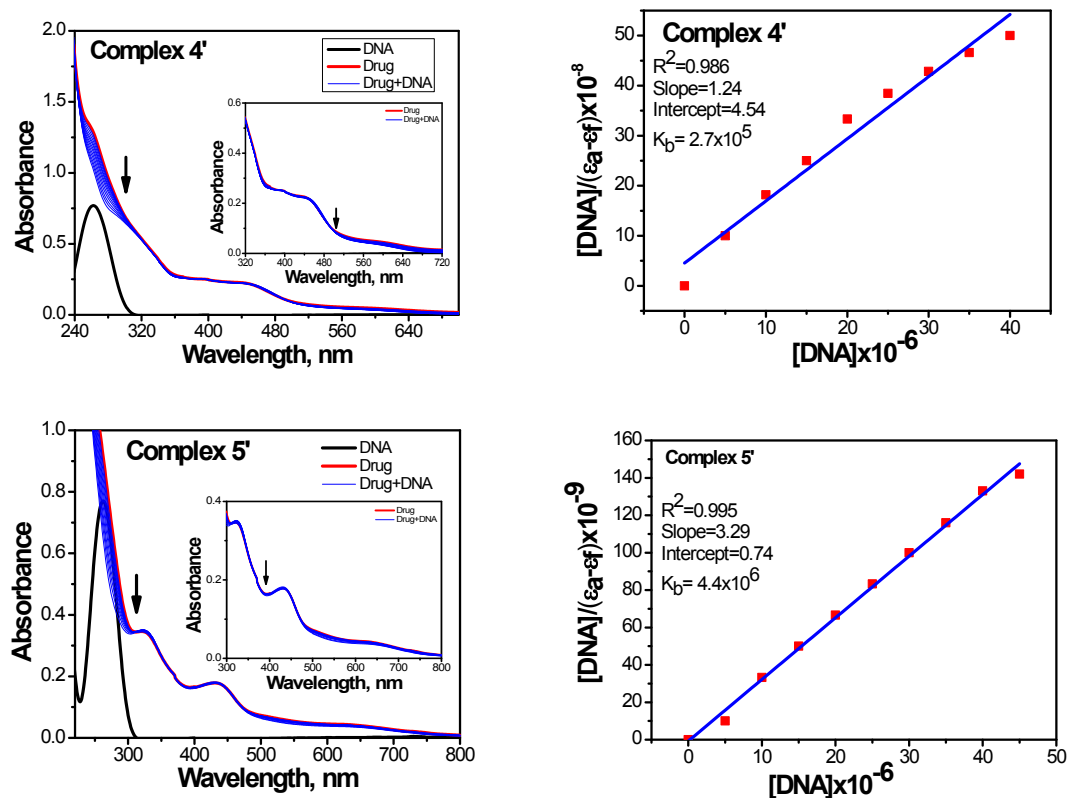
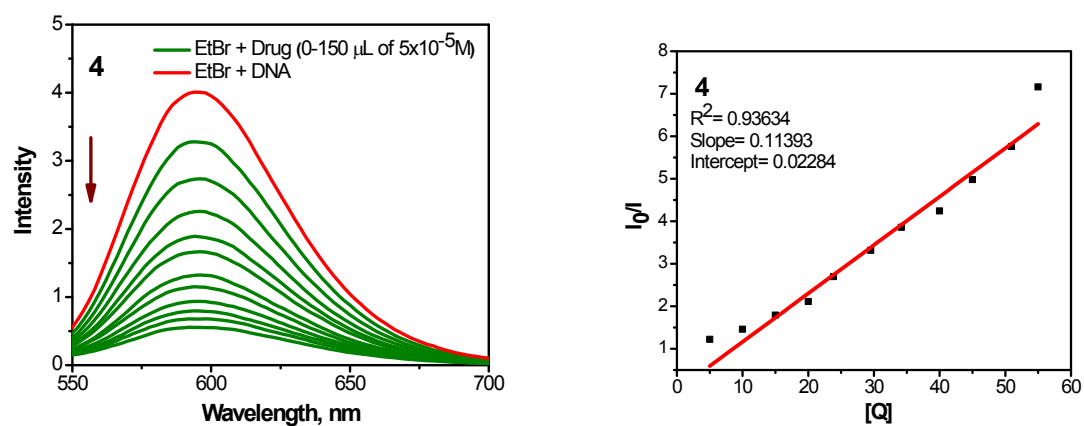


Fig. S5 (a) UV-visible titration of compounds **4**, **5**, **4'** and **5'** in 5 mM Tris-HCl/NaCl buffer with incremental addition of CT-DNA at pH 7.2. (b) $[DNA]/(\epsilon_a - \epsilon_f)$ vs. $[DNA]$ linear plots of complexes



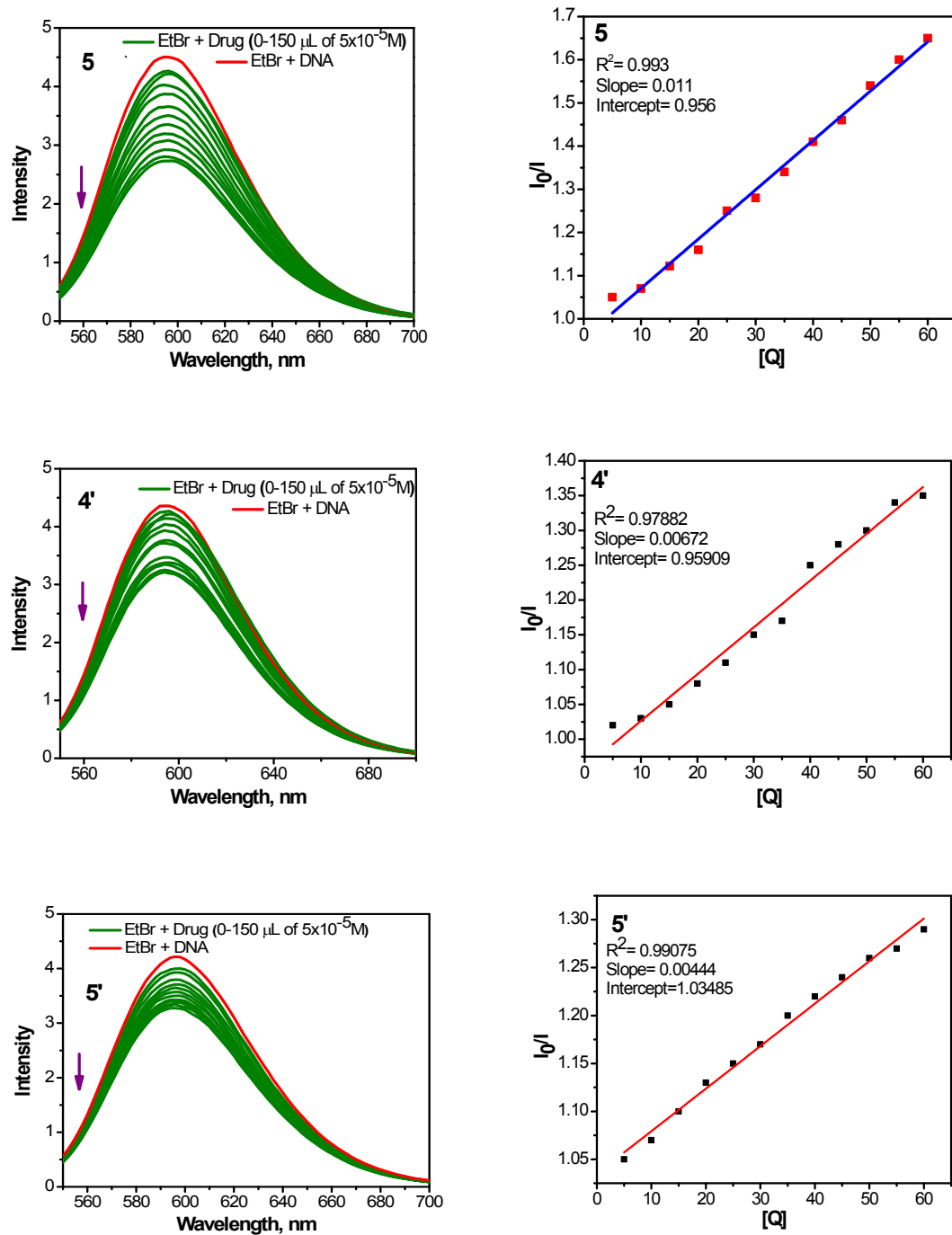
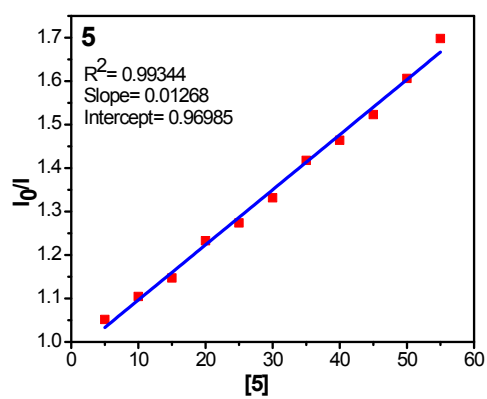
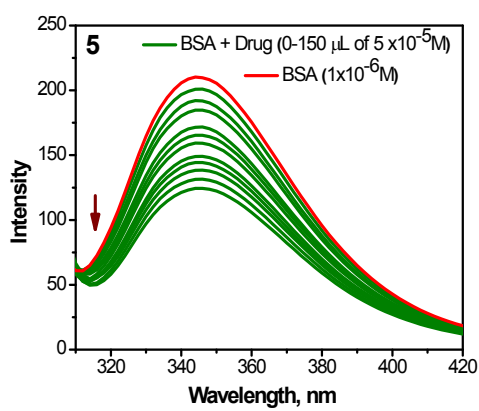
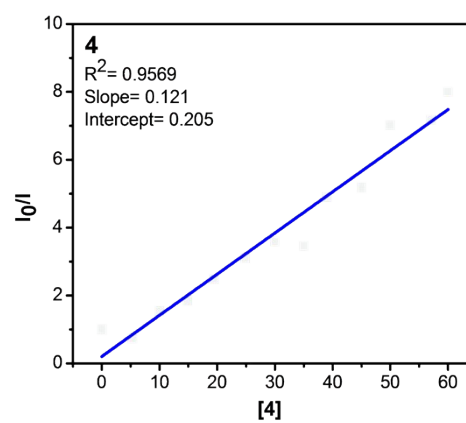
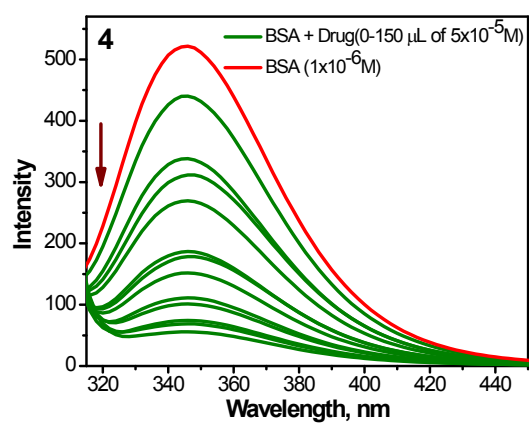


Fig. S6 Emission spectral traces of the ethidium bromide bound DNA with increasing concentration of compounds **4**, **5**, **4'** and **5'** in 5 mM Tris-HCl/NaCl buffer of pH 7.2. λ_{ex} = 485 nm, λ_{em} = 598 nm and 596 nm, [DNA] = 120 μ M, [EthB] = 8 μ M, [**4**]₅₀ = 30 μ M, [**4'**]₅₀ = 25 μ M, [**5**]₅₀ = 30 μ M, [**5'**]₅₀ = 25 μ M Stern-Volmer plots of I_0/I vs. Complex.



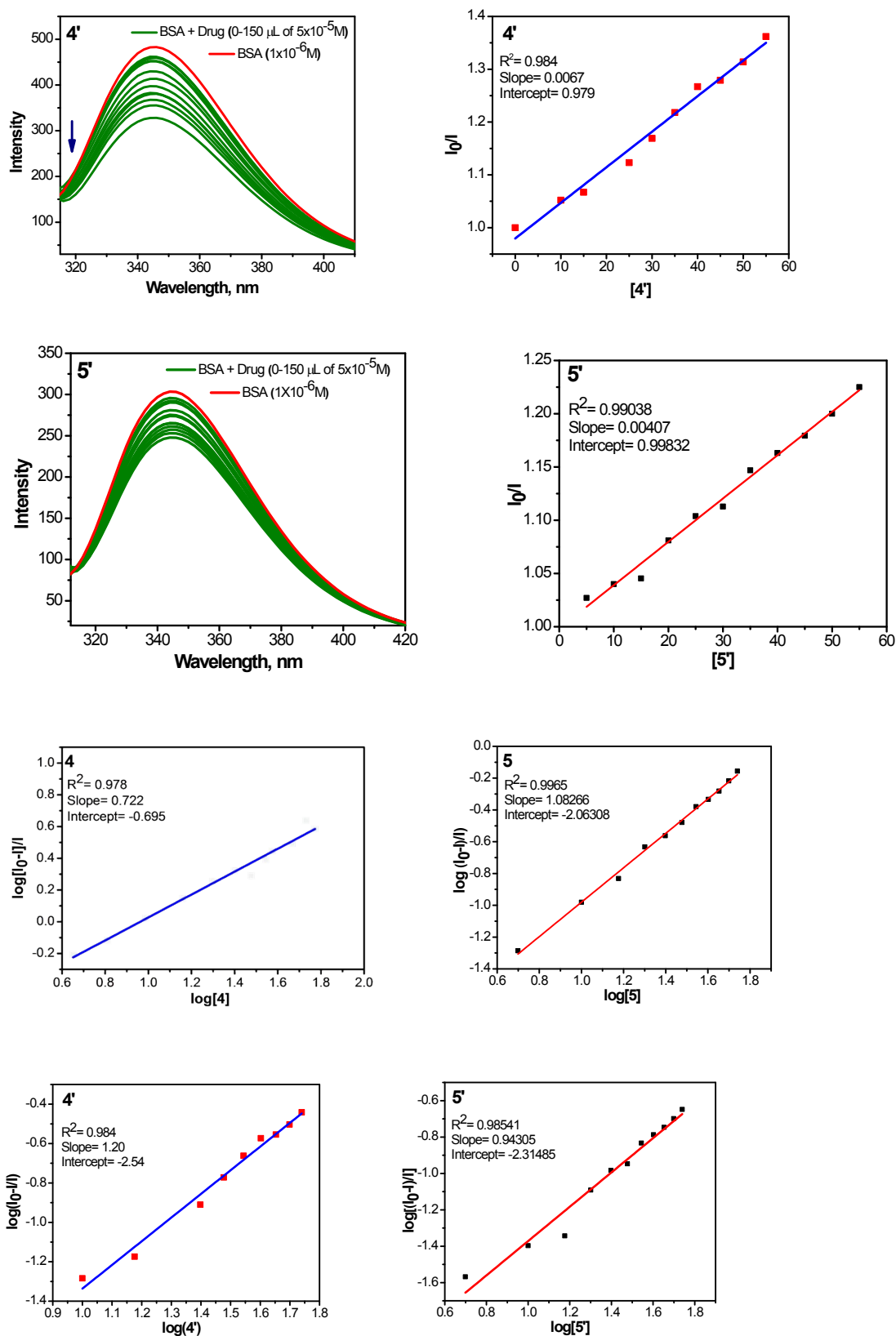


Fig. S7 (a) Fluorescence quenching of BSA on addition of compounds (a) **4**, **5**, **4'** and **5'** in 5 mM Tris HCl/NaCl buffer at pH 7.2 (λ_{ex} = 295; λ_{em} = 350 nm). (b) Plot of I_0/I vs. concentrations of compounds Scatchard plot of $\log ([I_0-I]/I)$ vs. $\log$ [compound] for BSA in the presence of complexes

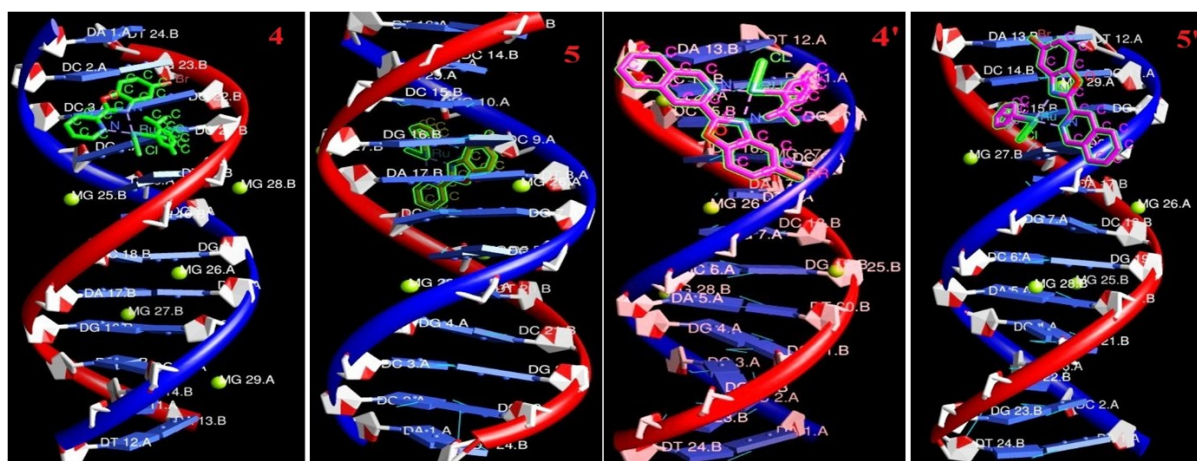


Fig. S8 Docking pose of compounds **4**, **5**, **4'** and **5'** within the minor groove of DNA

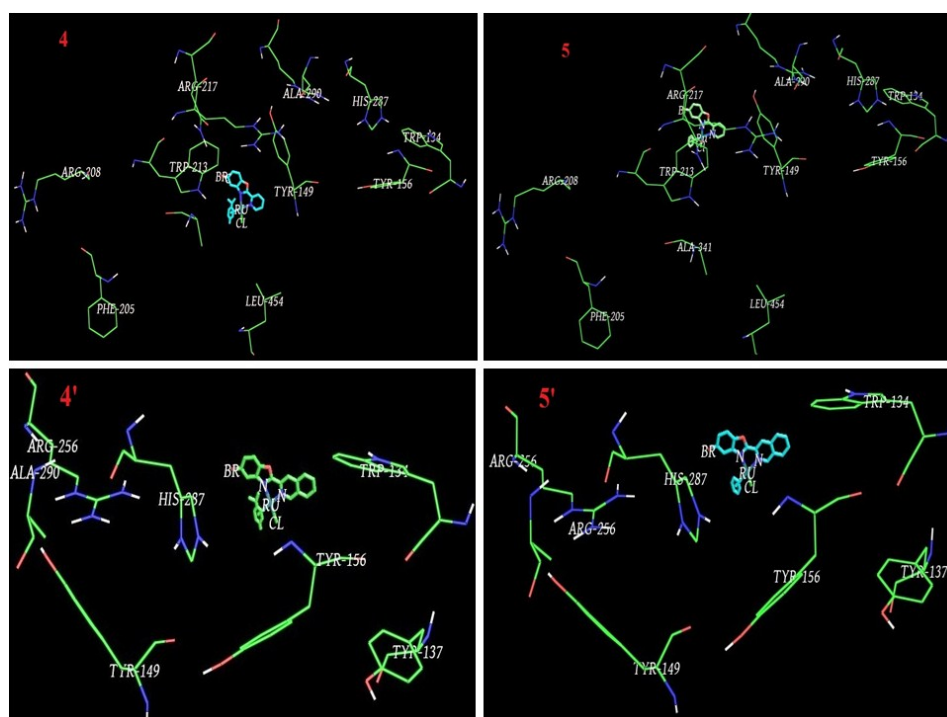


Fig. S9 Docking pose of compounds **4**, **5**, **4'** and **5'** within the active site of BSA protein

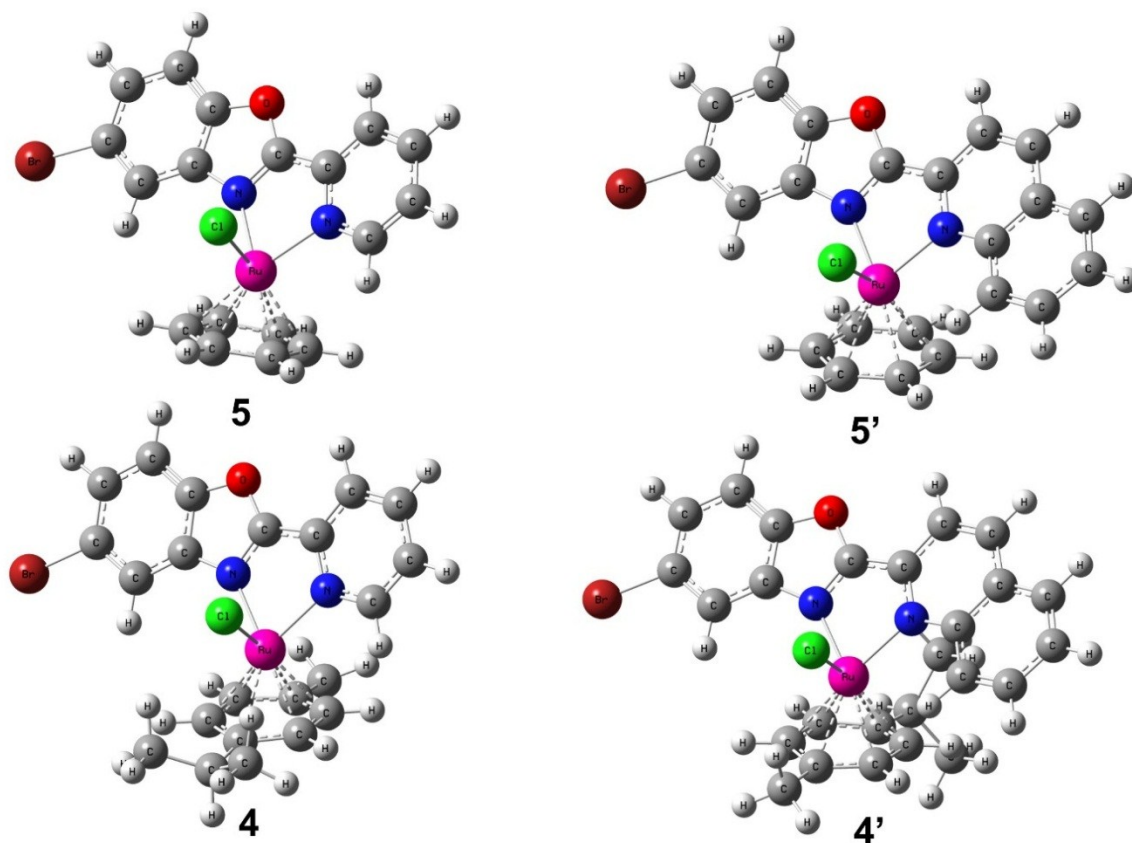


Fig. S10 Optimized structure of complexes (4, 5, 4' and 5')

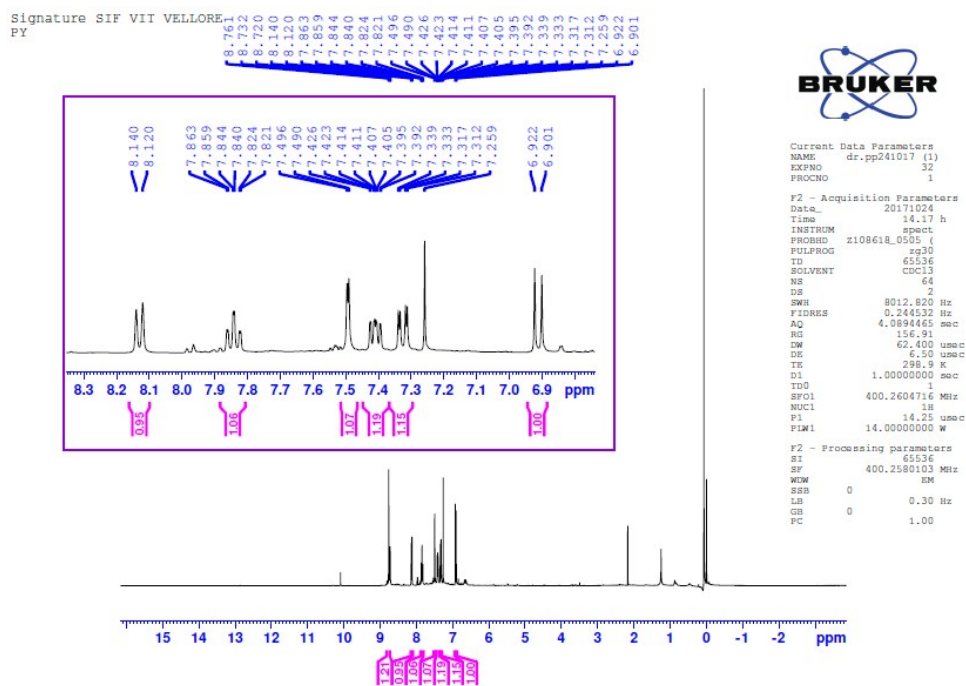


Fig. S11 ^1H NMR of compound 3

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PP

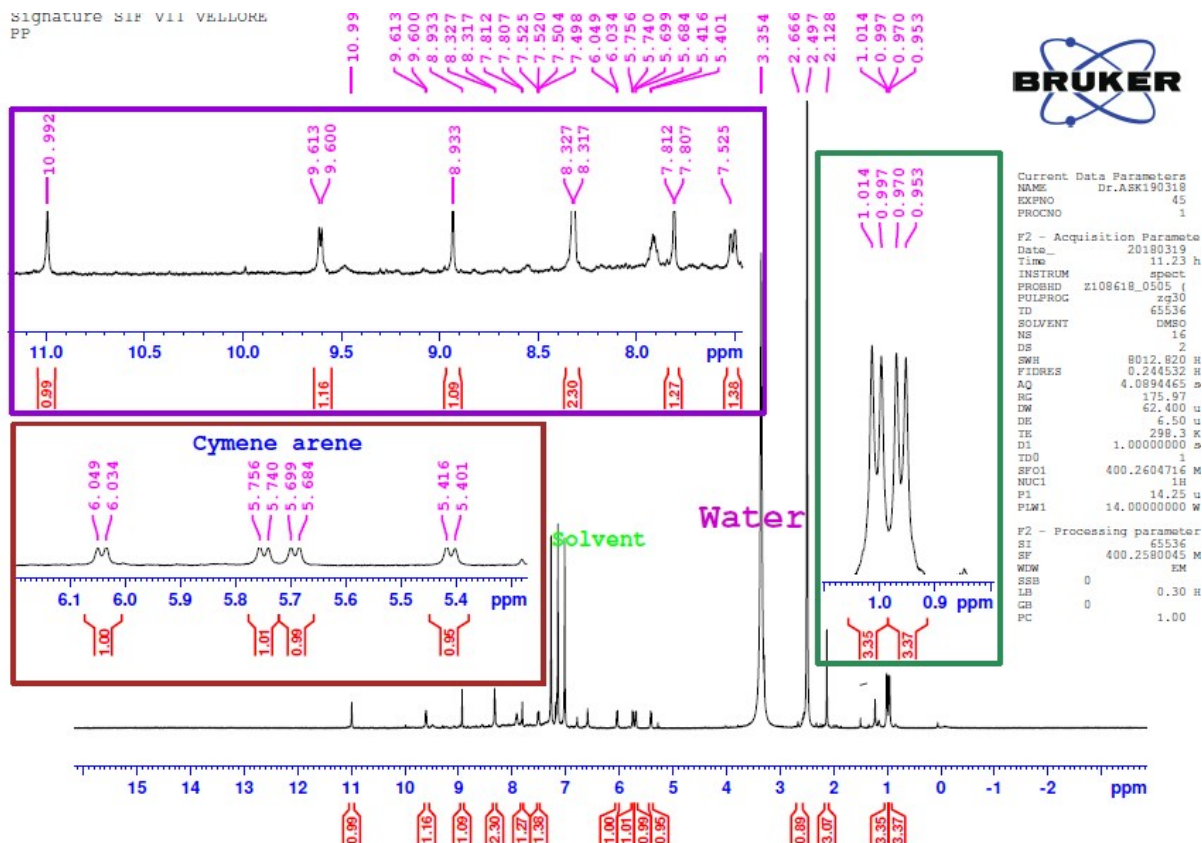


Fig. S12 ¹H NMR of complex 4

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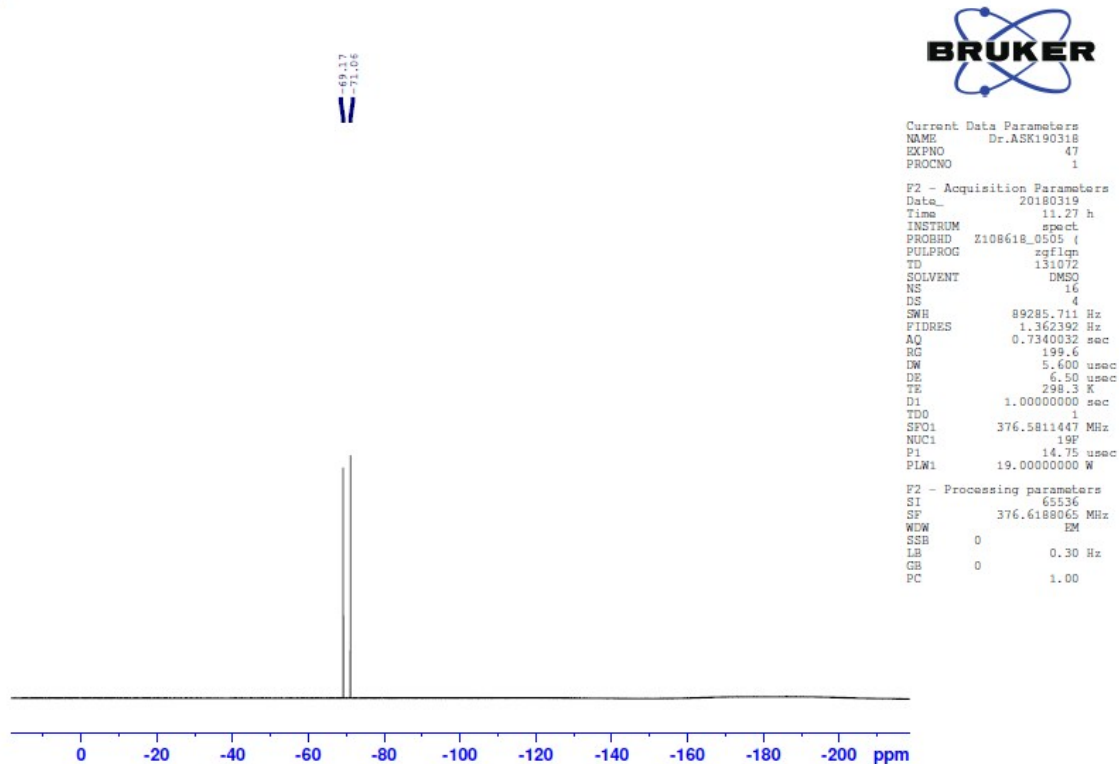
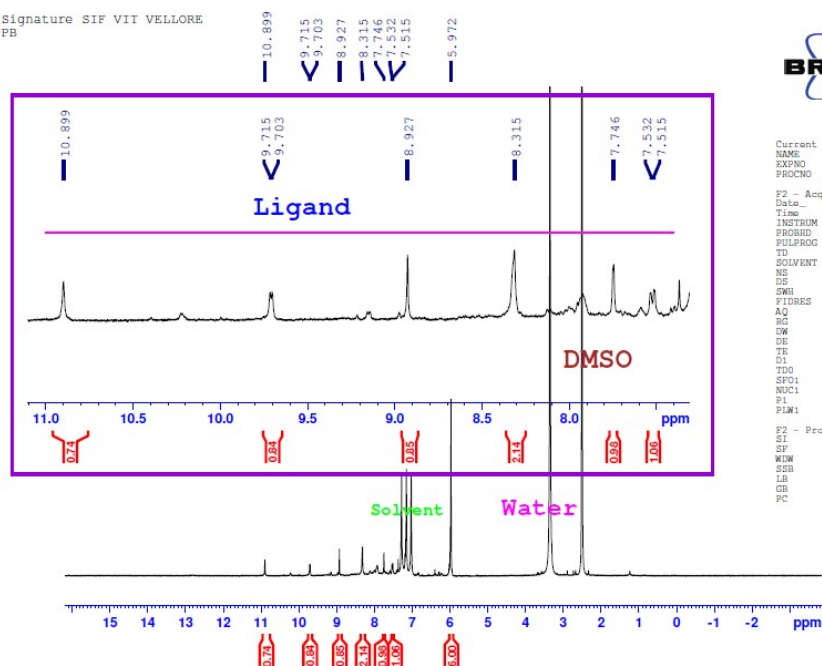


Fig. S13 ¹⁹F NMR of complex 4



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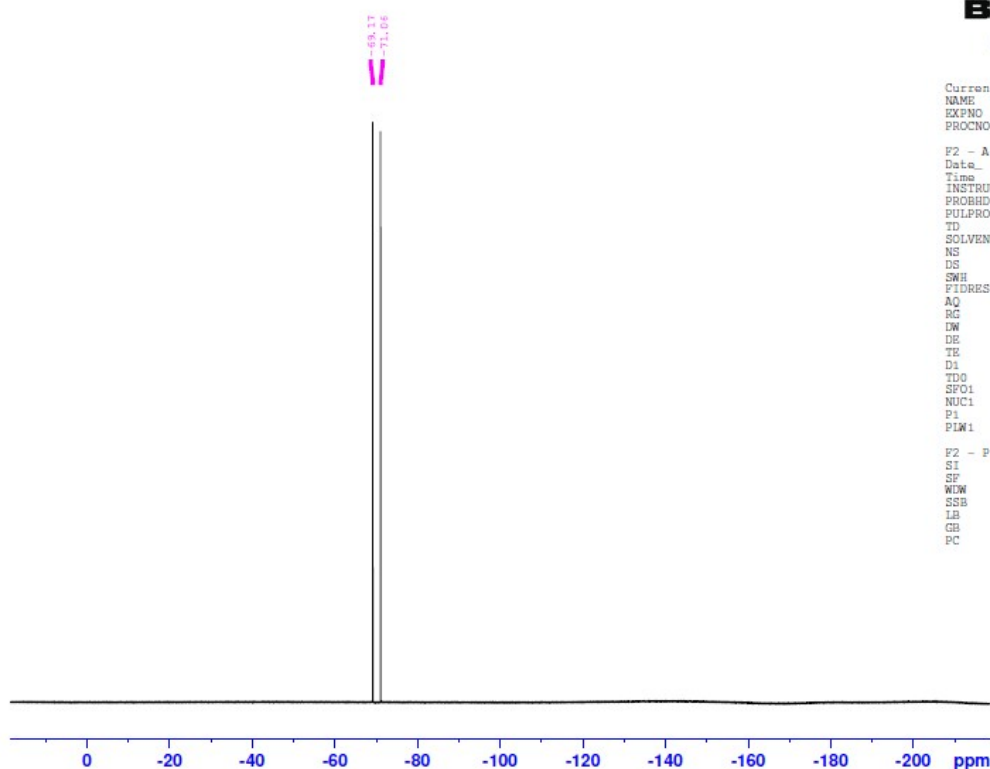
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Fig. S16 ¹H NMR spectra of complex 5

Signature SIF VII VELLORE
PB



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Fig. S17 ¹⁹F NMR spectra of complex 5

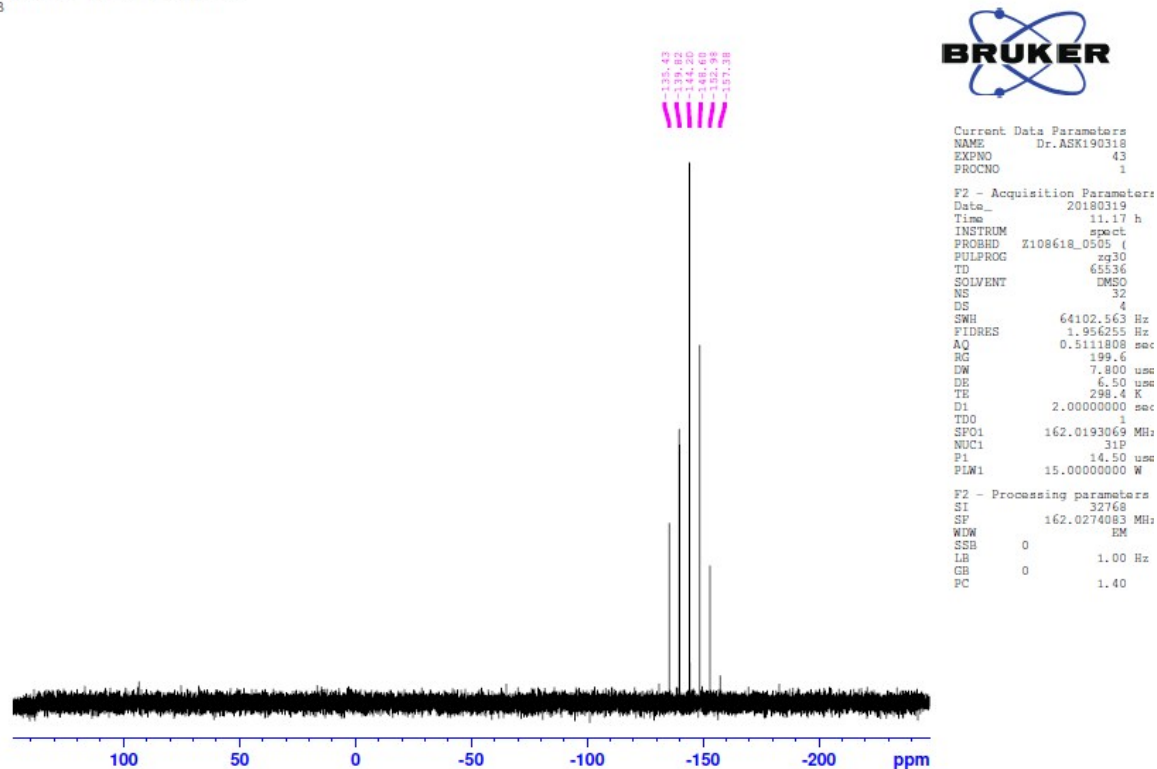


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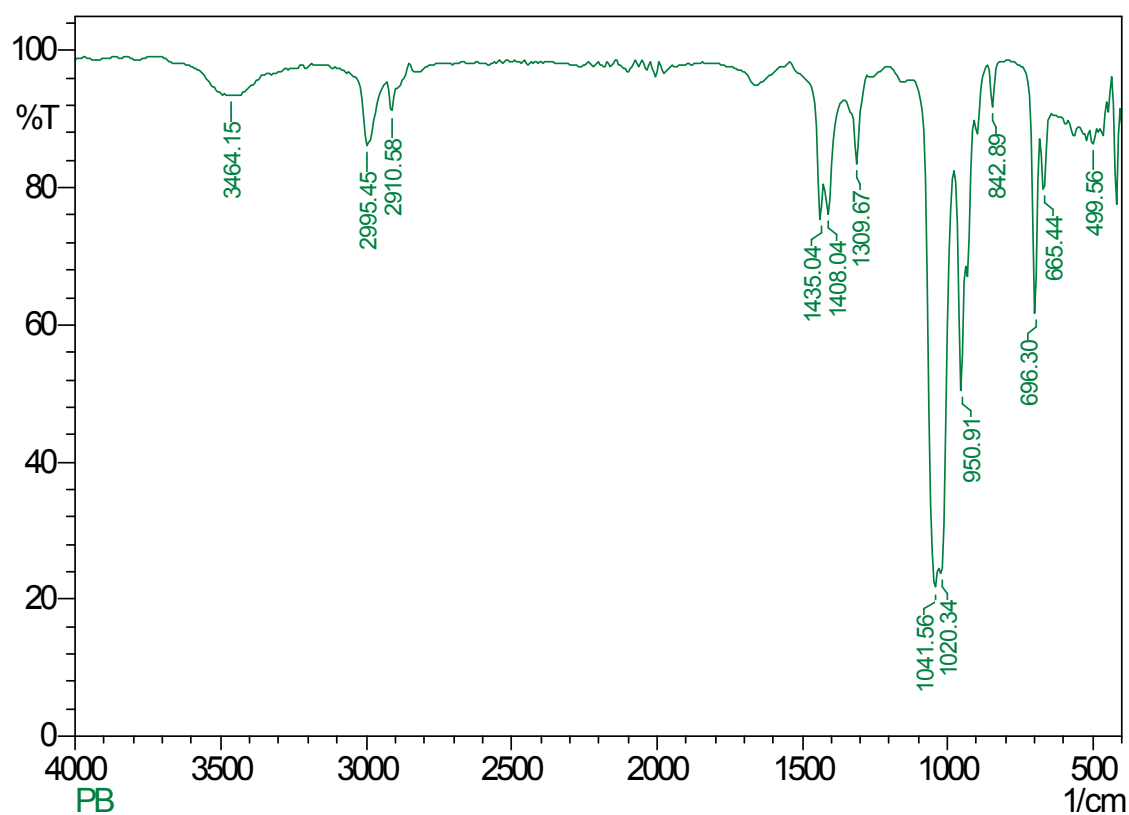


Fig. S19 FTIR spectra of complex 5

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Q.P

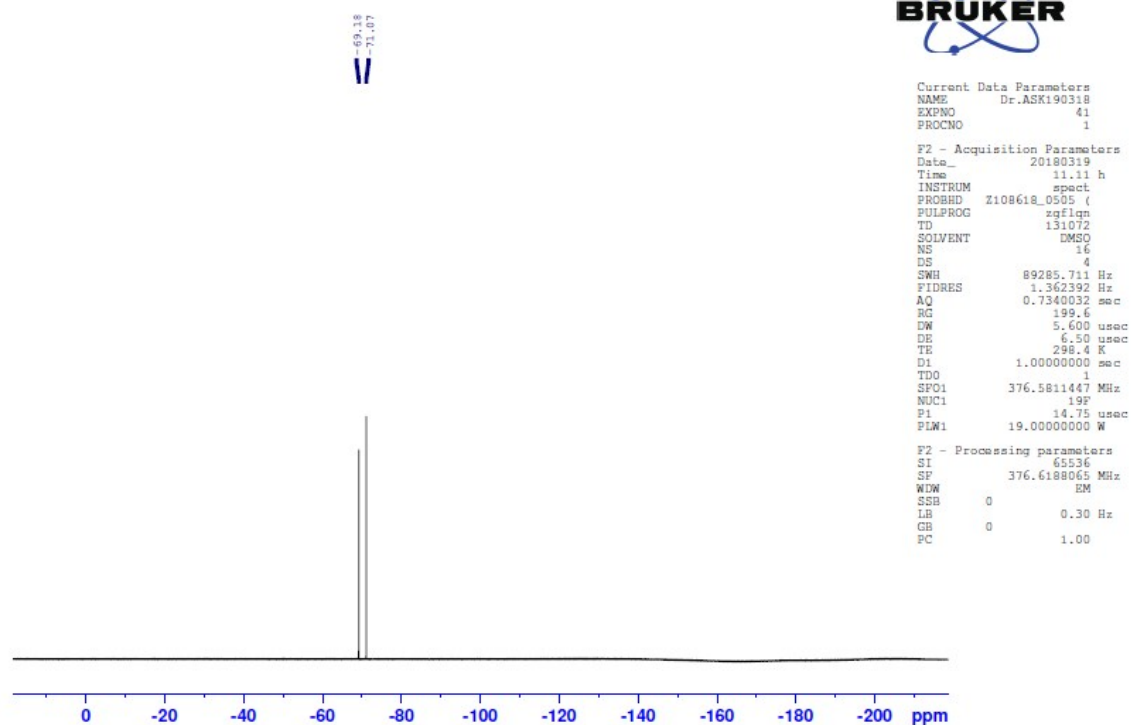


Fig. S22 ¹⁹F NMR spectra of complex 4'

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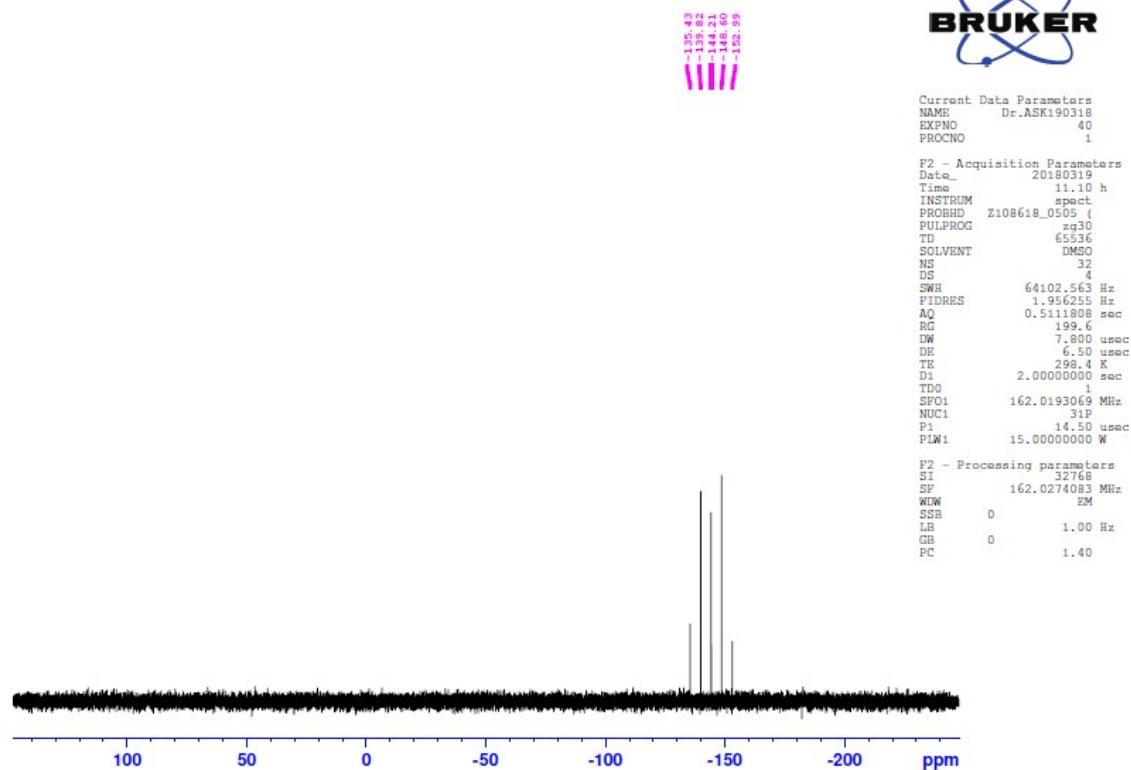


Fig. S23 ³¹P spectra of complex 4'

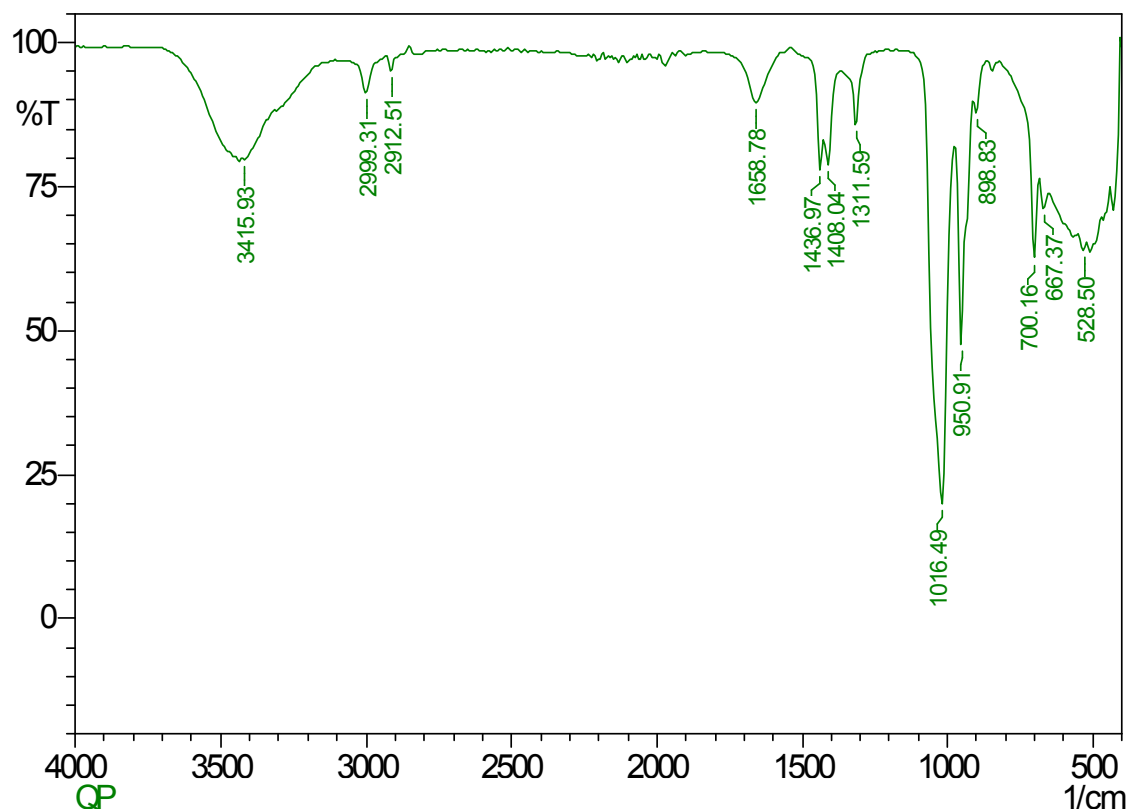


Fig. S24 FTIR spectra of complex 4'

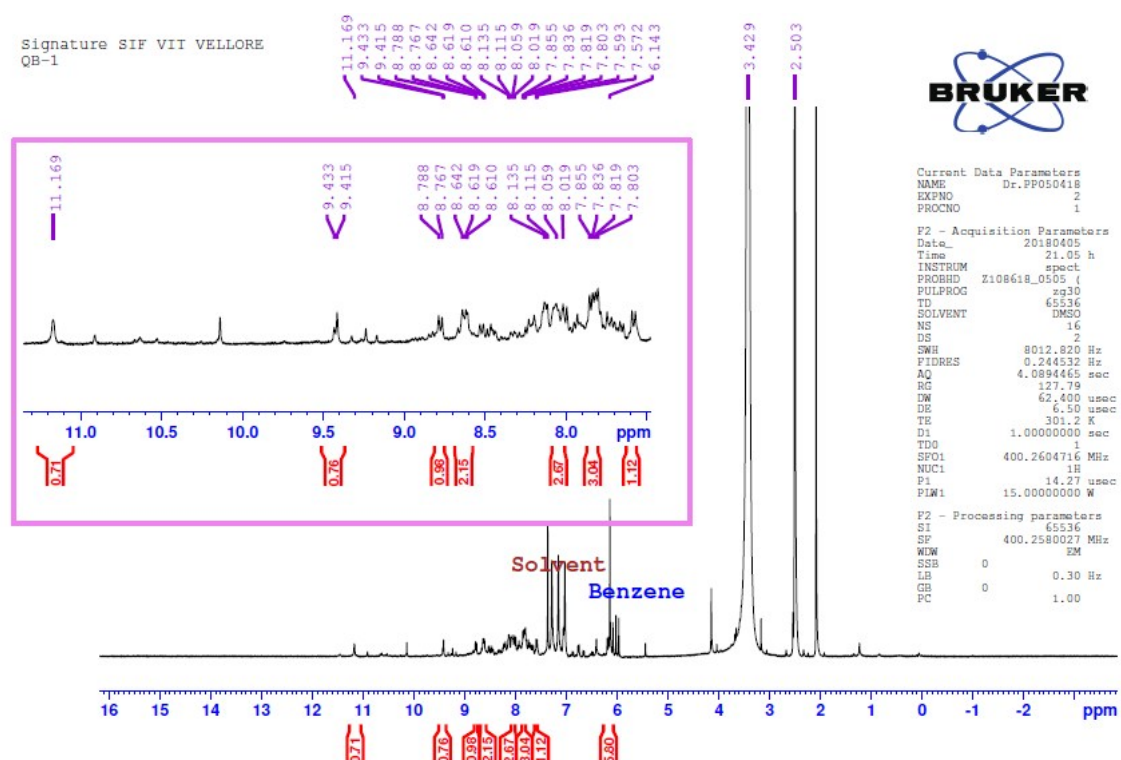
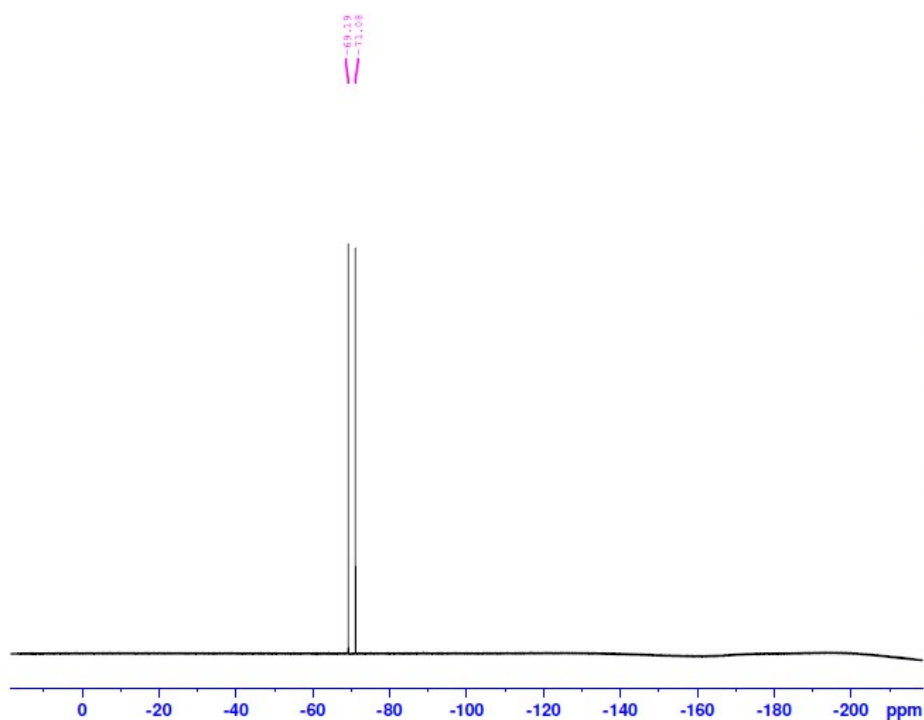


Fig. S25 ¹H spectra of complex 5'

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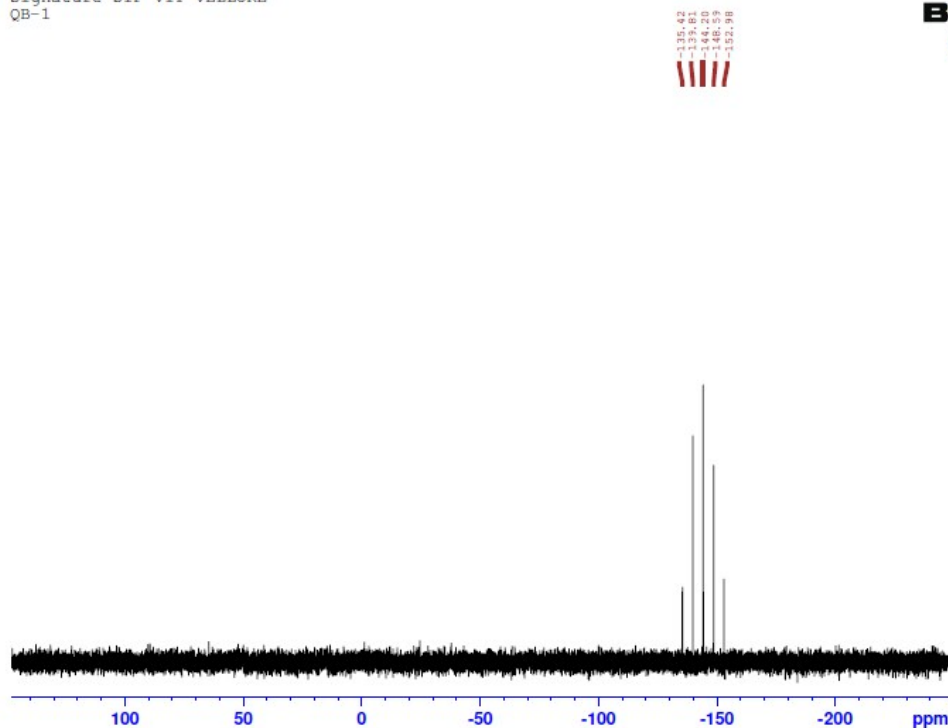
Current Data Parameters
NAME Dr.PP050418
EXPNO 3
PROCNO 1

F2 - Acquisition Parameters
Date_ 20180405
Time 21.07 h
INSTRUM spect
PROBHD Z108618_0505 ()
PULPROG zgpg30
TD 131072
SOLVENT DMSO
NS 16
DS 4
SWH 89285.711 Hz
FIDRES 1.362392 Hz
AQ 0.7340032 sec
RG 199.6
DW 5.600 usec
DE 6.50 usec
TE 301.2 K
D1 1.00000000 sec
TDO 1
SFO1 376.5811447 MHz
NUC1 19F
P1 14.75 usec
PLW1 19.00000000 W

F2 - Processing parameters
SI 65536
SF 376.6188065 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00

Fig. S26 ¹⁹F spectra of complex 5'

Signature SIF VIT VELLORE
QB-1



Current Data Parameters
NAME Dr.PP050418
EXPNO 4
PROCNO 1

F2 - Acquisition Parameters
Date_ 20180405
Time 21.10
INSTRUM spect
PROBHD Z108618_0505 ()
PULPROG zg30
TD 65536
SOLVENT DMSO
NS 32
DS 4
SWH 64102.563
FIDRES 1.956255
AQ 0.5111808
RG 199.6
DW 7.800
DE 6.50
TE 301.3
D1 2.00000000
TDO 1
SFO1 162.0193069 MHz
NUC1 31P
P1 14.50
PLW1 15.00000000 W

F2 - Processing parameters
SI 32768
SF 162.0274083 MHz
WDW EM
SSB 0
LB 1.00
GB 0
PC 1.40

Fig. S27 ³¹P spectra of complex 5'

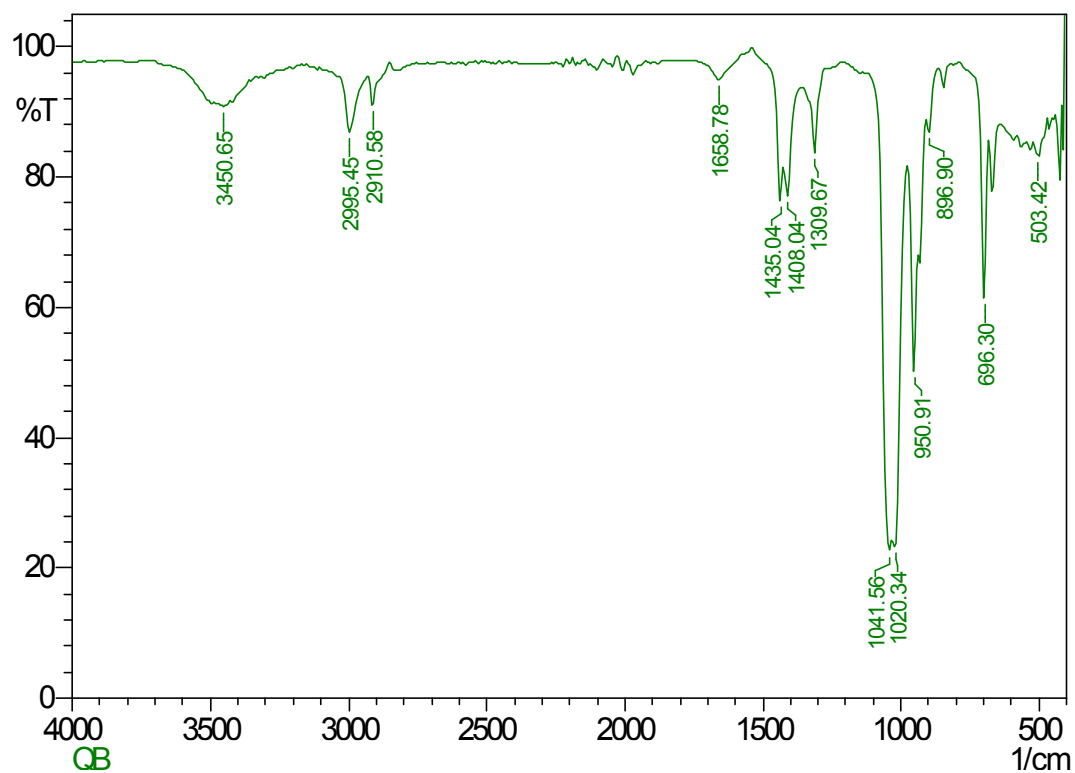


Fig. S28 FTIR spectra of complex 5'