

## **Electronic supplementary information**

### **Bimetallic Ru(II) and Os(II) complexes based on pyrene-bisimidazole spacer: synthesis, photophysics, electrochemistry and multisignalling DNA binding studies in the near infrared region**

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## Physical measurements

Elemental analyses of the compounds were performed with a Vario-Micro V2.0.11 elemental (CHNSO) analyzer. NMR spectra were collected on Bruker 500 spectrometer in DMSO-*d*<sub>6</sub>, and high resolution mass spectroscopy was performed on a Waters Xevo G2 QTOF mass spectrometer. The UV/vis absorption spectra were recorded with a Shimadzu UV 1800 spectrometer. A matched pair of quartz cuvettes (path length 1 cm) was employed. Steady state luminescence spectra were obtained either by a Perkin–Elmer LS55 or Spex fluorolog-2 spectrofluorometer equipped with DM3000F software. Luminescence quantum yields were determined by using literature method taking [Ru(bpy)<sub>3</sub>]<sup>2+</sup> as the standard. Luminescence lifetime measurements were carried out by using time–correlated single photon counting set up from Horiba Jobin-Yvon. The luminescence decay data were collected on a Hamamatsu MCP photomultiplier (R3809) and were analyzed by using IBH DAS6 software. Cyclic and square-wave voltammetric experiments were performed in deaerated acetonitrile with a BAS epsilon electrochemistry system and a three-electrode set up consisting of a platinum or glassy carbon working electrode, a platinum counter electrode, and Ag/AgCl reference electrode. Tetraethylammonium perchlorate (TEAP) was used as background electrolyte. The potentials reported in this study were referenced against the Ag/AgCl electrode, which under the given experimental conditions gave a value of 0.36 V for the Fc/Fc<sup>+</sup> couple. Spectroelectrochemical measurements were performed with a system consisting of a BAS epsilon potentiostat/galvanostat, a Shimadzu 3600 UV-VIS-NIR spectrophotometer and an optically transparent spectroelectrochemical cell specially designed by BAS.

## X-ray crystallographic analyses

The crystallographic data, details of data collection, and refinement parameter for the complex **1** are summarized in Table S1. Single crystal of suitable size was obtained by diffusing toluene to 1:1 acetonitrile-dichloromethane solution of the complex. The crystal was immersed in paratone oil and then mounted on the tip of a glass fibre and cemented using epoxy resin. Intensity data for the crystal was collected using MoK $\alpha$  ( $\lambda = 0.7107 \text{ \AA}$ ) radiation on a Bruker SMART APEX II diffractometer equipped with CCD area detector at room temperature. The data integration and reduction were processed with SAINT<sup>S1</sup> software provided with the software package of SMART APEX II. An empirical absorption correction was applied to the collected reflections with SADABS.<sup>S1</sup> The structure was solved by direct methods using SHELXTL<sup>S2</sup> and was refined on F<sup>2</sup> by the full-matrix least-squares technique

using the SHELXL-97<sup>S3</sup> program package. Graphics was generated using PLATON.<sup>S4</sup> Non-hydrogen atoms were refined anisotropically until the convergence. All the hydrogen atoms were geometrically positioned and treated as riding atoms.

### Theoretical computational methods

Quantum chemical calculations were performed with the Gaussian 09 program<sup>S5</sup> employing the DFT method with Becke's three-parameter hybrid functional and Lee-Yang-Parr's gradient corrected correlation functional B3LYP level of theory.<sup>S6,S7</sup> The 6-31G(d) basis set was employed for the C, H and N while SDD basis set was used for Ru and Os atoms.<sup>[S8]</sup> Geometries were fully optimized using the criteria of the respective programs. Time-dependent DFT (TD-DFT) calculations were performed to assign the experimentally observed bands in the complexes.<sup>S9-S12</sup> The excitation energies, computed within the acetonitrile solvent simulated by the CPCM model,<sup>S13</sup> has been determined by using the so-called nonequilibrium approach, which has been designed for the study of the absorption process.<sup>S14,S15</sup> Only singlet-singlet transitions, that is, the spin-allowed transitions, have been taken into account. UKS calculations were also performed to calculate singlet-triplet energy gap in CH<sub>3</sub>CN using the CPCM model. Orbital analysis was completed with Gauss View<sup>S16</sup> and Gauss sum 2.2.<sup>S17</sup>

### DNA binding experiments

#### Absorption and emission spectral experiments

UV-vis absorption and emission titrations were carried out by maintaining a constant metal complex concentration (20 μM) and adding incrementally the CT-DNA solution of appropriate concentration in 5 mM Tris-HCl/NaCl buffer with the ionic strength of 50 .0 × 10<sup>-3</sup> m (pH 7.30) at room temperature. After each addition of DNA to the solution of the metal complexes, the resulting solution was allowed to equilibrate at 25 °C for 5 min, after which the absorption and emission readings were noted. Proper corrections were made to the absorbance of CT-DNA. The equilibrium binding constant (K<sub>b</sub>) and the binding site size (s, per base pair) of the complexes to CT-DNA were calculated by non-linear least square analysis of the isotherm using the expression of Bard and co-workers based on the McGhee-von Hippel (MvH) model.<sup>S18-S19</sup>

$$(\epsilon_a - \epsilon_f)/(\epsilon_b - \epsilon_f) = (b - (b^2 - 2K_b^2C_t[\text{DNA}]/s)^{1/2})/2K_bC_t \quad (\text{S1})$$

$$(I_a - I_f)/(I_b - I_f) = (b - (b^2 - 2K_b^2C_t[\text{DNA}]/s)^{1/2})/2K_bC_t \quad (\text{S2})$$

where  $b = 1 + K_b C_t + K_b [\text{DNA}]/2s$ ,  $\epsilon_a$  is the extinction coefficient observed for the spectral band at a given DNA concentration,  $\epsilon_f$  is the extinction coefficient of the free complex in solution,  $\epsilon_b$  is the extinction coefficient of the complex when fully bound to DNA. Similarly  $I_a$ ,  $I_f$  and  $I_b$  are the luminescence intensities of the complex at a given DNA concentration, free complex and fully bound to DNA respectively.  $K_b$  is the equilibrium binding constant,  $C_t$  is the total complex concentration,  $[\text{DNA}]$  is the DNA concentration in nucleotides and  $s$  is the binding site size of the complexes in base pairs. The non-linear least-square fit analysis was done using Origin Lab software. The lifetimes of the complexes were also recorded as a function of CT-DNA in 5 mM Tris-HCl/ NaCl buffer with the ionic strength of  $50.0 \times 10^{-3}$  M (pH 7.30) at room temperature.

### Competitive binding fluorescence experiments

The apparent binding constants ( $K_{app}$ ) of the complexes have been evaluated by using ethidium bromide (EB)-bound CT-DNA solution in 5 mM Tris-HCl/ NaCl buffer with the ionic strength of  $50.0 \times 10^{-3}$  M (pH 7.30) at room temperature. The luminescence intensities at 602 nm (546 nm excitation) of EB-DNA conjugate with increasing concentration of **1** and **2** were recorded. The  $K_{app}$  values were calculated by using the equation  $K_{EB} \times [\text{EB}] = K_{app} \times [\text{complex}]$ , where  $K_{EB}$  is the binding constant of EB ( $K_{EB} = 1.25 \times 10^6 \text{ M}^{-1}$ ),  $[\text{EB}]$  is the concentration of EB (13  $\mu\text{M}$ ) and  $[\text{complex}]$  is the concentration of the complex at 50% reduction of the initial fluorescence emission intensity.<sup>S20</sup>

### Circular dichroism experiments

The Circular dichroism (CD) spectroscopy was studied on a JASCO J-815 CD spectropolarimeter between 400 and 200 nm in continuous scanning mode at 25°C. Experiments were performed by adding progressively increasing amounts of complexes to the solutions of CT-DNA [40 $\mu\text{M}$ ] in 5 mM Tris-HCl/ NaCl buffer with the ionic strength of  $50.0 \times 10^{-3}$  M (pH 7.30).

### Thermal denaturation experiments

The effect of temperature on CT-DNA in absence and presence of the complexes were measured from circular dichroism study. The experiments were performed on a JASCO J-815 CD spectropolarimeter equipped with a Peltier temperature-controller (JASCO-PTC-423S/15). CD melting profiles were generated by rising temperature from 40 to 110°C with

monitoring the absorbance at 260nm in 5 mM Tris-HCl/ NaCl buffer with the ionic strength of  $50.0 \times 10^{-3}$  M (pH 7.30) medium. The melting temperature  $T_m$  was taken as the midpoint of the melting curve which was obtained from the maximum of the first derivative plot and checked graphically by tangent method.  $\Delta T_m$  value was calculated by subtracting  $T_m$  of the free nucleic acid from that of the complex.<sup>S21</sup>

### **Molecular docking studies**

Molecular docking studies were performed using HEX 8.0 software.<sup>S22</sup> To perform such studies, crystal structure of B-DNA (PDB ID: 1BNA) and DFT optimized structures (using Gaussian 09 software) of the complexes were used. The crystal structure of B-DNA was retrieved from the protein data bank (<http://www.rcsb.org./pdb>). To obtain the molecular docked structures and to elucidate different type of interaction processes between DNA and the receptors Biovia Discovery Studio 2016 software was used.

### **Viscometric study**

The viscosity of sonicated DNA<sup>S23</sup> was measured by a fabricated micro viscometer, maintained at  $28 (\pm 0.5)$  °C in a thermostatic water bath. Then the viscosities of CT-DNA-EB, CT-DNA-Complex **1** and CTDNA-Complex **2** conjugates were measured in the same experimental conditions. Data were presented as  $(\eta/\eta_0)^{1/3}$  versus the ratio of the concentration of either EB or complex **1** or **2** to that of the CT DNA, where  $\eta_0$  is the viscosity of CT DNA solution alone and  $\eta$  is the viscosity of CT DNA solution in the presence of either the complexes or EB. Viscosity values were calculated from the observed flow time of CT DNA by the relation  $\eta = t - t_0$ , where  $t$  and  $t_0$  are the values of flow times for the solution and the Tris-NaCl buffer respectively.

**Table S1** Crystallographic data for complex **1**

Compound	<b>1</b>
Formula	C75 H56 Cl4 N14 O16 Ru2
fw	1753.28
T (K)	293(2)
Cryst. Syst.	Triclinic
Space group	P-1
<i>a</i> (Å)	12.2111(5)
<i>b</i> (Å)	18.6407(6)
<i>c</i> (Å)	18.8598(8)
$\alpha$ (deg)	75.740(3)
$\beta$ (deg)	77.216(4)
$\gamma$ (deg)	80.547(3)
<i>V</i> (Å <sup>3</sup> )	4030.1(3)
Dc(g cm <sup>-3</sup> )	1.445
<i>Z</i>	2
$\mu$ (mm <sup>-1</sup> )	0.580
<i>F</i> (000)	1776.0
$\theta$ range (deg)	1.72-25.00
Data/restraints/params	14064/0/1001
GOF on <i>F</i> <sup>2</sup>	1.003
<i>R</i> <sub>1</sub> [ <i>I</i> > 2σ( <i>I</i> )] <sup>a</sup> ,	0.0656
<i>wR</i> <sub>2</sub> (all data) <sup>b</sup>	0.1423
Δρ <sub>max</sub> /Δρ <sub>min</sub> (e Å <sup>-3</sup> )	1.846/ -1.533
<sup>a</sup> R1( <i>F</i> ) = [ $\sum    F_0   -  F_C    / \sum   F_0  $ ], <sup>b</sup> wR2 ( <i>F</i> <sup>2</sup> ) = [ $\sum w(F_0^2 - F_C^2)^2 / \sum w(F_0^2)^2$ ] <sup>1/2</sup>	

**Table S2** Selected calculated bond distances (Å) for **1** and **2** in ground and UKS optimized state along with available X-ray crystal data

	Exp <b>1</b>	Soln. <b>1</b>	UKS <b>1</b>			Soln. <b>2</b>	UKS <b>2</b>
Ru1-N1	2.08(5)	2.11	2.12		Os1-N1	2.11	2.12
Ru1-N2	2.14(4)	2.21	2.21		Os1-N2	2.20	2.20
Ru1-N8	2.04(5)	2.10	2.09		Os1-N8	2.10	2.10
Ru1-N7	2.06(5)	2.10	2.13		Os1-N7	2.10	2.13
Ru1-N9	2.04(5)	2.09	2.06		Os1-N9	2.09	2.08
Ru1-N10	2.04(4)	2.07	2.05		Os1-N10	2.08	2.07
Ru2-N4	2.16(4)	2.21	2.21		Os2-N4	2.20	2.20
Ru2-N6	2.10(4)	2.11	2.11		Os2-N6	2.12	2.12
Ru2-N14	2.07(5)	2.10	2.10		Os2-N14	2.10	2.10
Ru2-N13	2.05(5)	2.10	2.10		Os2-N13	2.10	2.10
Ru2-N12	2.05(5)	2.09	2.09		Os2-N12	2.09	2.09
Ru2-N11	2.04(4)	2.07	2.07		Os2-N11	2.08	2.08

**Table S3** Selected calculated bond angles (°) for **1** and **2** in ground and UKS optimized state along with available X-ray crystal data

	<b>1</b>				<b>2</b>		
	Exp <b>1</b>	Soln. <b>1</b>	UKS <b>1</b>			Soln. <b>2</b>	UKS <b>2</b>
N1-Ru1-N2	78.20(17)	77.82	77.60		N1-Os1-N2	77.10	77.11
N1-Ru1-N8	172.33(19)	173.69	173.73		N1- Os1-N8	173.39	173.87
N1-Ru1-N7	93.74(19)	96.33	98.14		N1- Os1-N7	96.69	98.69
N1-Ru1-N9	92.11(18)	88.43	86.36		N1- Os1-N9	88.14	86.36
N1-Ru1-N10	94.88(19)	93.98	93.33		N1- Os1-N10	94.50	93.93
N2-Ru1-N8	100.25(17)	103.81	105.87		N2- Os1-N8	104.56	106.28
N2-Ru1-N7	87.82(17)	86.54	84.46		N2- Os1-N7	85.95	84.34
N2-Ru1-N9	96.79(18)	97.50	98.37		N2- Os1-N9	98.16	98.57
N2-Ru1-N10	172.08(18)	171.01	170.85		N2- Os1-N10	170.85	170.88
N8-Ru1-N7	78.66(19)	77.78	77.20		N8- Os1-N7	77.14	76.78
N8-Ru1-N9	95.54(19)	97.32	98.14		N8- Os1-N9	97.88	98.03
N8-Ru1-N10	87.12(19)	84.72	83.25		N8- Os1-N10	84.19	82.78
N7-Ru1-N9	173.19(19)	174.32	175.11		N7- Os1-N9	174.26	174.63
N7-Ru1-N10	96.54(18)	98.12	98.21		N7- Os1-N10	98.77	98.85
N9-Ru1-N10	79.48(18)	78.41	79.59		N9- Os1-N10	77.70	78.93
N4-Ru2-N12	169.38(18)	170.96	170.86		N4-Os2-N12	170.75	170.77
N4-Ru2-N14	103.08(17)	103.89	104.00		N4- Os2-N14	104.66	104.64
N4-Ru2-N11	94.86(18)	97.46	97.31		N4- Os2-N11	98.08	98.00
N4-Ru2-N13	90.08(18)	86.56	86.64		N4- Os2-N13	86.01	86.04
N6-Ru2-N4	78.44(17)	77.81	77.79		N6- Os2-N4	77.10	77.08
N6-Ru2-N14	176.46(18)	173.70	173.62		N6- Os2-N14	173.41	173.29
N6-Ru2-N13	97.38(18)	96.36	96.30		N6- Os2-N13	96.74	96.62
N6-Ru2-N11	88.25(18)	88.43	88.45		N6- Os2-N11	88.14	88.19
N6-Ru2-N12	92.12(18)	93.94	93.91		N6- Os2-N12	94.42	94.49
N13-Ru2-N11	173.17(19)	174.13	174.37		N13- Os2-N11	174.23	174.31
N13-Ru2-N12	96.01(19)	98.15	98.21		N13- Os2-N12	98.80	98.83
N11-Ru2-N12	79.85(19)	78.41	78.42		N11- Os2-N12	77.70	77.71
N14-Ru2-N13	79.47(18)	77.78	77.78		N14- Os2-N13	77.15	77.14
N14-Ru2-N11	94.78(18)	97.29	97.31		N14- Os2-N11	97.81	97.89
N14-Ru2-N12	86.63(18)	84.69	84.64		N14- Os2-N12	84.16	84.14



**Table S4** Selected molecular orbital along with their energies and compositions for **1** and **2** in solution phase

	Energy/ eV	(%) Composition			
		<b>1 (%) Composition</b>			
MO	<b>1</b>	Ru <sup>II</sup>	pyrene imida	pyridine	bpy
LUMO+3	-2.59	5.33	12.02	12.49	70.14
LUMO+2	-2.61	6.27	3.52	1.06	89.13
LUMO+1	-2.65	2.06	9.13	7.10	81.68
LUMO	-2.65	1.81	6.68	5.48	86.02
HOMO	-5.93	23.83	65.59	7.25	3.30
HOMO-1	-6.09	80.08	5.59	3.44	10.87
HOMO-2	-6.18	60.22	26.06	5.06	8.64
HOMO-3	-6.27	73.44	10.11	3.52	12.91
		<b>2 (%) Composition</b>			
		Os <sup>II</sup>	pyrene imida	pyridine	bpy
MO	<b>2</b>	8.02	12.25	12.99	66.73
LUMO+3	-2.58	8.42	11.97	6.42	73.17
LUMO+2	-2.59	1.60	13.26	15.97	69.15
LUMO+1	-2.71	1.63	10.42	12.96	74.96
LUMO	-2.72	65.64	19.23	4.64	10.47
HOMO	-5.77	79.23	4.54	3.75	12.46
HOMO-1	-5.81	70.77	8.67	4.58	15.96
HOMO-2	-6.04	67.74	10.08	5.49	16.67
HOMO-3	-6.05				

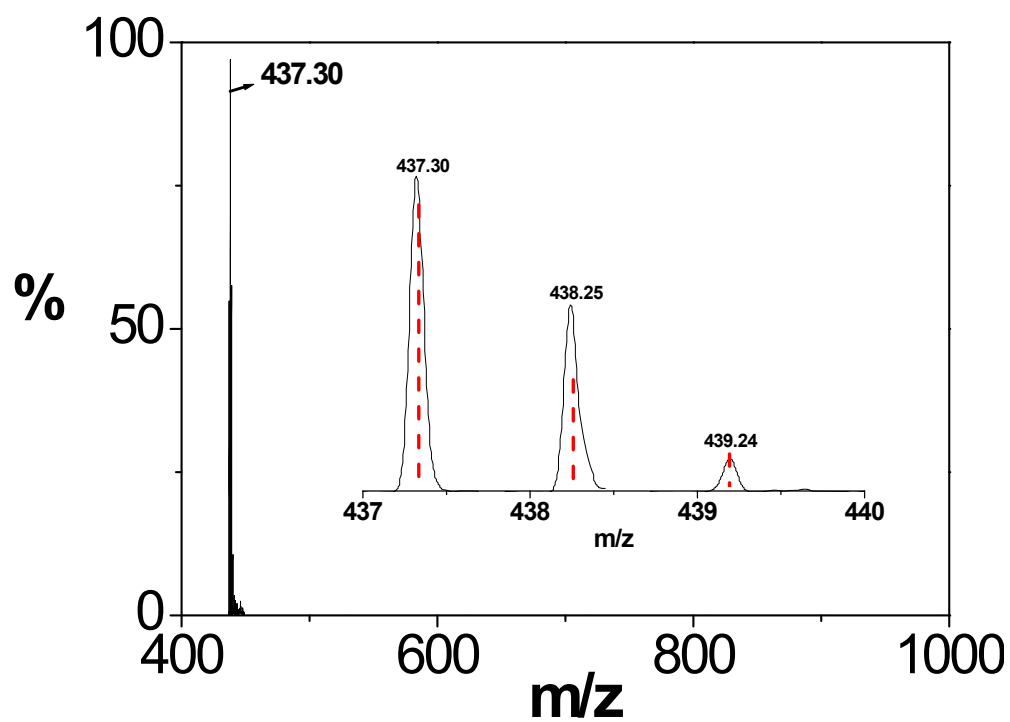
**Table S5** Selected molecular orbital along with their energies and compositions for **1** and **2** in UKS optimized state

	Energy/ eV	(%) Composition			
		<b>1 (%) Composition</b>			
MO	<b>1</b>	Ru <sup>II</sup>	pyrene imida	pyridine	bpy
LUMO+3	-2.61	5.94	4.11	4.86	85.07
LUMO+2	-2.66	1.66	6.09	9.03	83.21
LUMO+1	-2.86	2.37	38.44	57.61	1.55
LUMO	-2.95	2.90	0.51	0.15	96.42
HOMO	-3.89	3.81	10.91	0.51	84.75
HOMO-1	-6.02	38.25	49.99	6.45	5.29
HOMO-2	-6.19	44.93	42.48	6.02	6.54
HOMO-3	-6.30	75.12	8.95	2.53	13.39
		<b>2 (%) Composition</b>			
		Os <sup>II</sup>	pyrene imida	pyridine	bpy
MO	<b>2</b>	4.31	20.57	26.06	49.03
LUMO+3	-2.64	4.54	31.17	45.31	18.97
LUMO+2	-2.89	5.60	3.85	5.09	85.45
LUMO+1	-2.94	3.50	7.61	4.67	84.02
LUMO	-3.15	3.80	8.45	4.01	83.71
HOMO	-3.38	53.86	30.14	5.43	10.54
HOMO-1	-6.00				

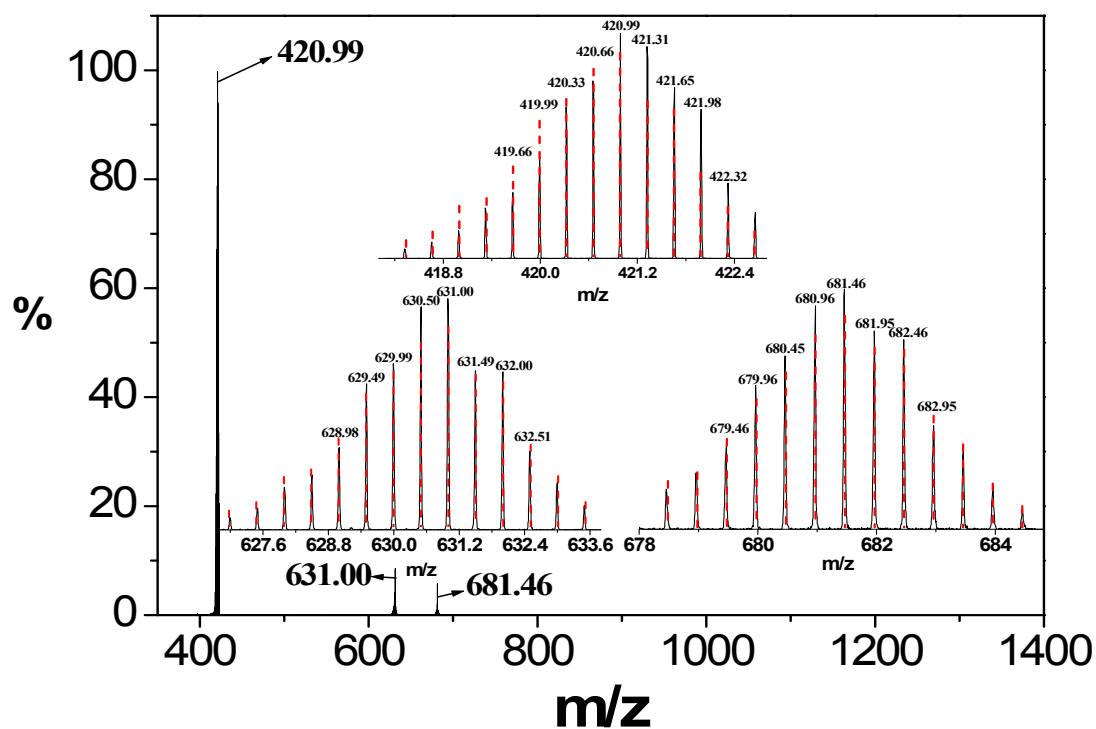
HOMO-2	-6.16		67.05	11.44	5.54	15.96
HOMO-3	-6.20		31.92	58.09	4.54	5.42

**Table S6** Binding constants of **1** and **2** with DNA in 50mM, 300mM and 1000mM ionic strength tris buffer solution (pH=7.30)

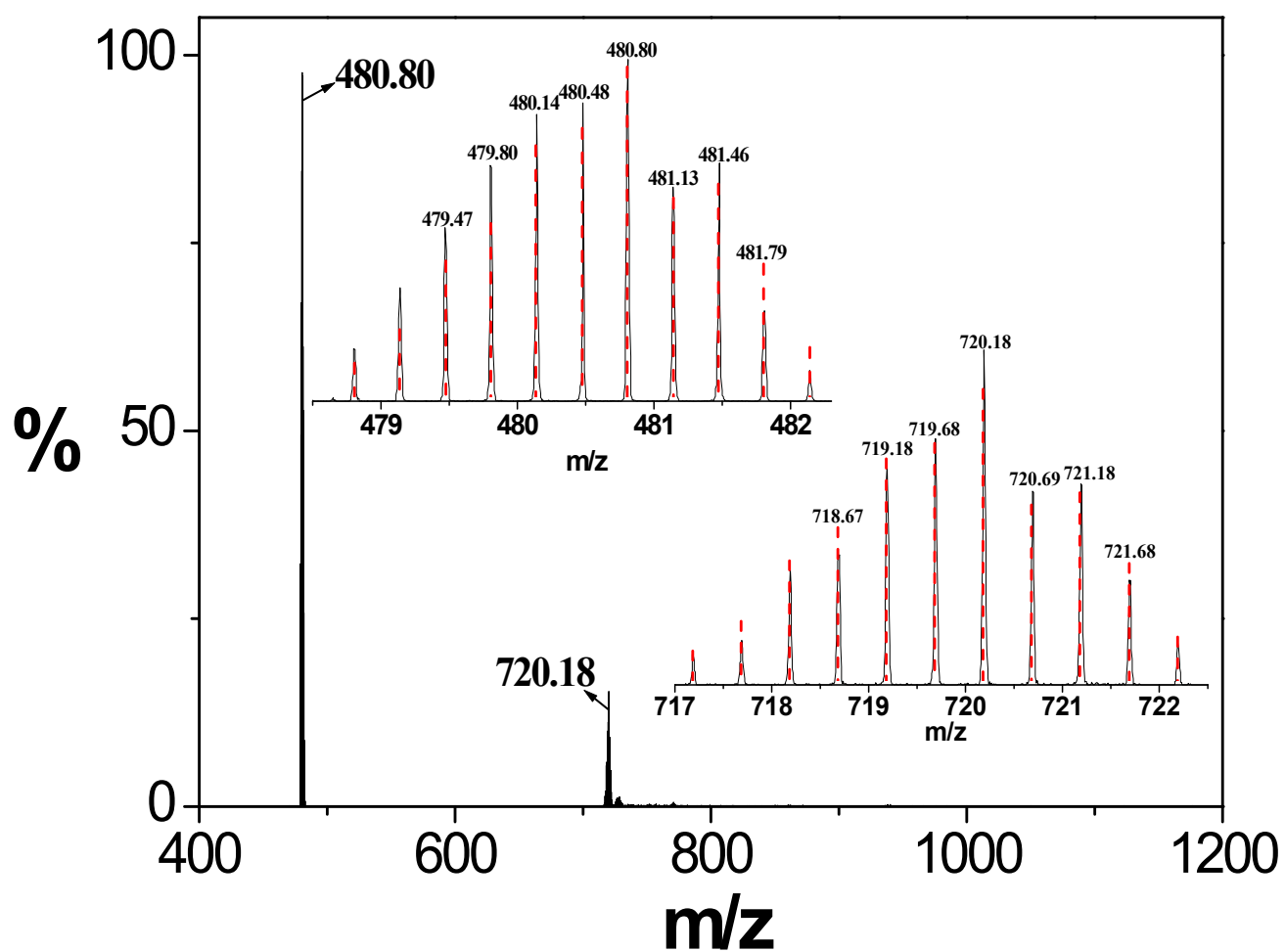
Complex	From Absorption( $K_b$ )	From Emission( $K_b$ )
<b>Strength 50 mM</b>		
<b>1</b>	$K_b = (7.23 \pm 0.59) \times 10^6 \text{ M}^{-1}$ $s = 2.07 \pm 0.02$	$K_b = (7.04 \pm 0.65) \times 10^6 \text{ M}^{-1}$ $s = 1.98 \pm 0.02$
<b>2</b>	$K_b = (7.51 \pm 0.65) \times 10^6 \text{ M}^{-1}$ $s = 1.96 \pm 0.02$	$K_b = 7.74 \pm 0.64 \times 10^6 \text{ M}^{-1}$ $s = 1.99 \pm 0.01$
<b>Strength 300 mM</b>		
<b>1</b>	$K_b = (7.62 \pm 0.17) \times 10^5 \text{ M}^{-1}$ $s = 2.04 \pm 0.05$	$K_b = (7.72 \pm 0.14) \times 10^5 \text{ M}^{-1}$ $s = 1.99 \pm 0.02$
<b>2</b>	$K_b = (8.14 \pm 0.14) \times 10^5 \text{ M}^{-1}$ $s = 2.06 \pm 0.03$	$K_b = (8.04 \pm 0.08) \times 10^5 \text{ M}^{-1}$ $s = 1.98 \pm 0.05$
<b>Strength 1000 mM</b>		
<b>1</b>	$K_b = (3.14 \pm 0.21) \times 10^5 \text{ M}^{-1}$ $s = 1.99 \pm 0.08$	$K_b = (3.32 \pm 0.19) \times 10^5 \text{ M}^{-1}$ $s = 2.03 \pm 0.04$
<b>2</b>	$K_b = (4.24 \pm 0.17) \times 10^5 \text{ M}^{-1}$ $s = 2.09 \pm 0.02$	$K_b = (4.76 \pm 0.11) \times 10^5 \text{ M}^{-1}$ $s = 2.10 \pm 0.02$



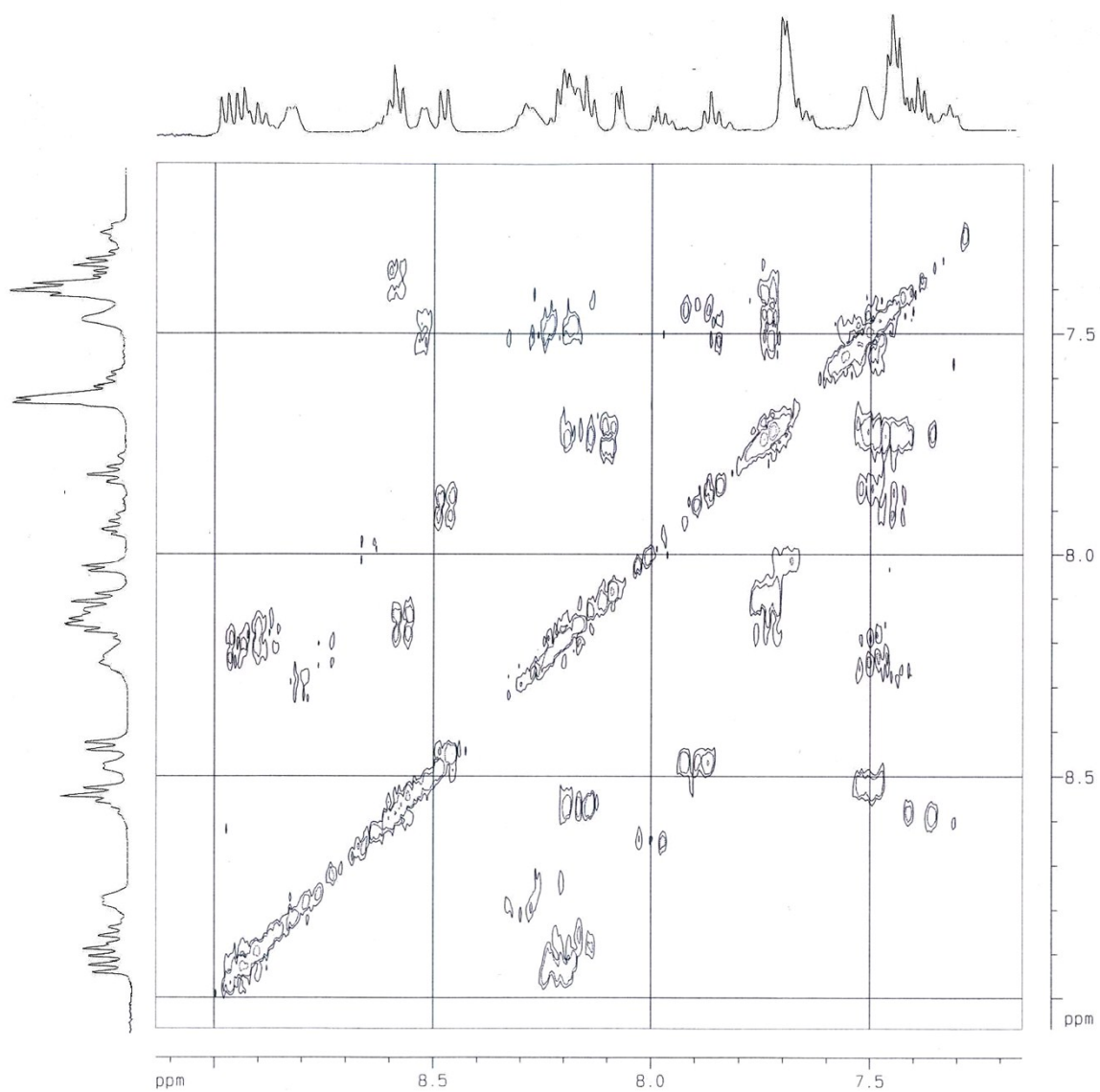
**Fig. S1** ESI-MS (positive) for the cation  $\text{H}_3\text{Imz}_2\text{PPy}_2^+$  ( $m/z = 437.30$ ) in acetonitrile showing the observed and isotopic distribution patterns.



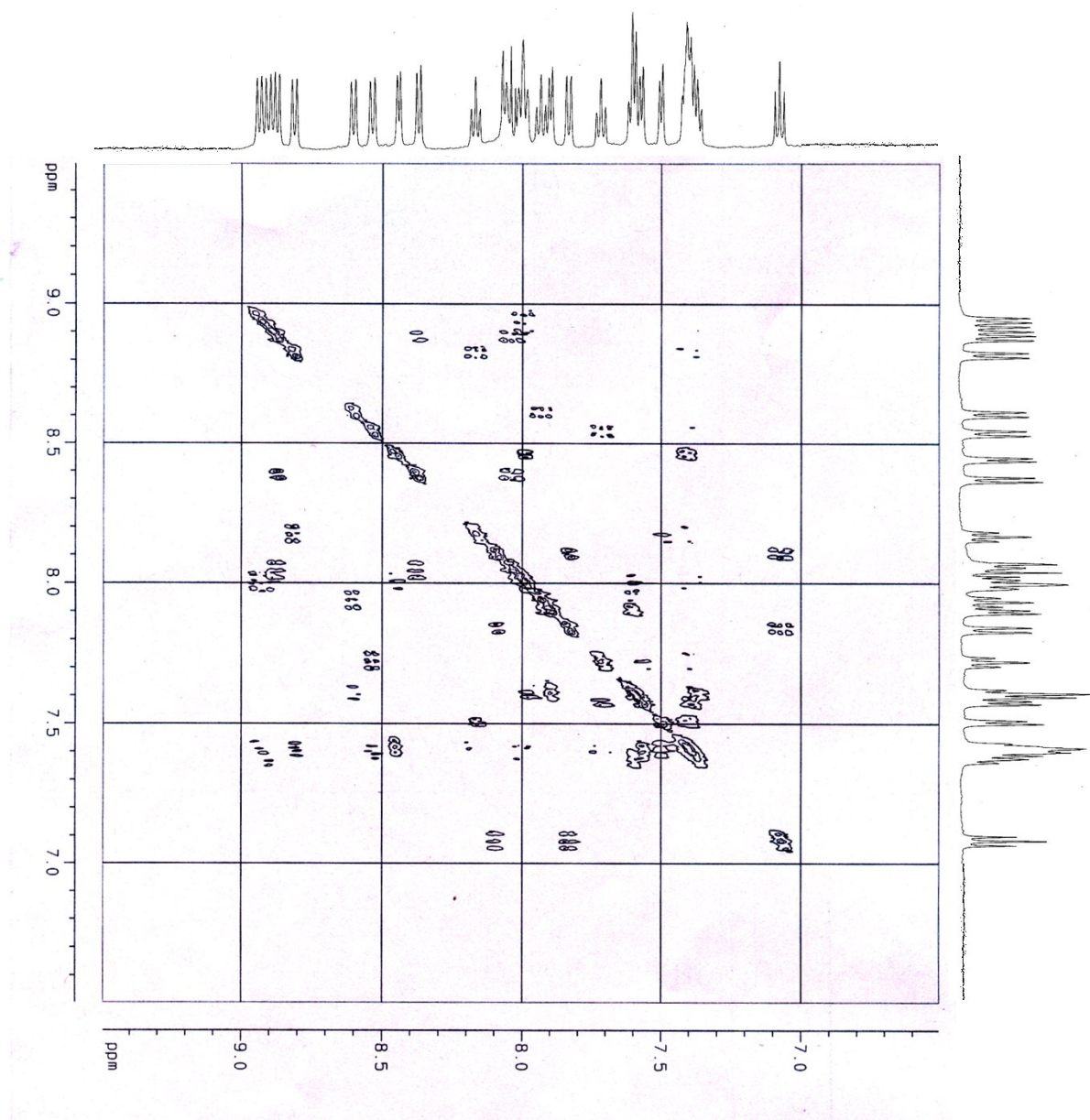
**Fig. S2** ESI-MS (positive) for the complex cations of **1**, [(bpy)<sub>2</sub>Ru(HImz<sub>2</sub>PPy<sub>2</sub>)Ru(bpy)<sub>2</sub>]<sup>3+</sup> (*m/z* = 420.99), [(bpy)<sub>2</sub>Ru(Imz<sub>2</sub>PPy<sub>2</sub>)Ru(bpy)<sub>2</sub>]<sup>2+</sup> (*m/z* = 631.00) and [(bpy)<sub>2</sub>Ru(HImz<sub>2</sub>PPy<sub>2</sub>)Ru(bpy)<sub>2</sub>(ClO<sub>4</sub>)]<sup>2+</sup> (*m/z* = 681.46) in acetonitrile showing both experimental and simulated isotopic distribution patterns.



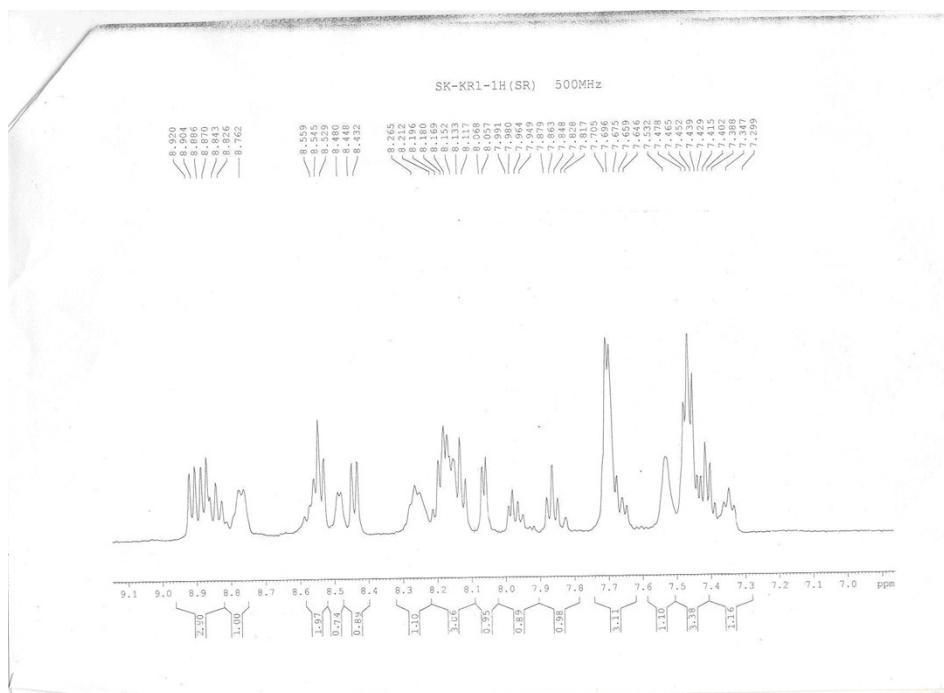
**Fig. S3** ESI-MS (positive) for the complex cations of **2**,  $[(bpy)_2Os(HImz_2PPy_2)Os(bpy)_2]^{3+}$  ( $m/z = 480.80$ ),  $[(bpy)_2Os(Imz_2PPy_2)Os(bpy)_2]^{2+}$  ( $m/z = 720.18$ ) in acetonitrile showing the observed and isotopic distribution patterns.



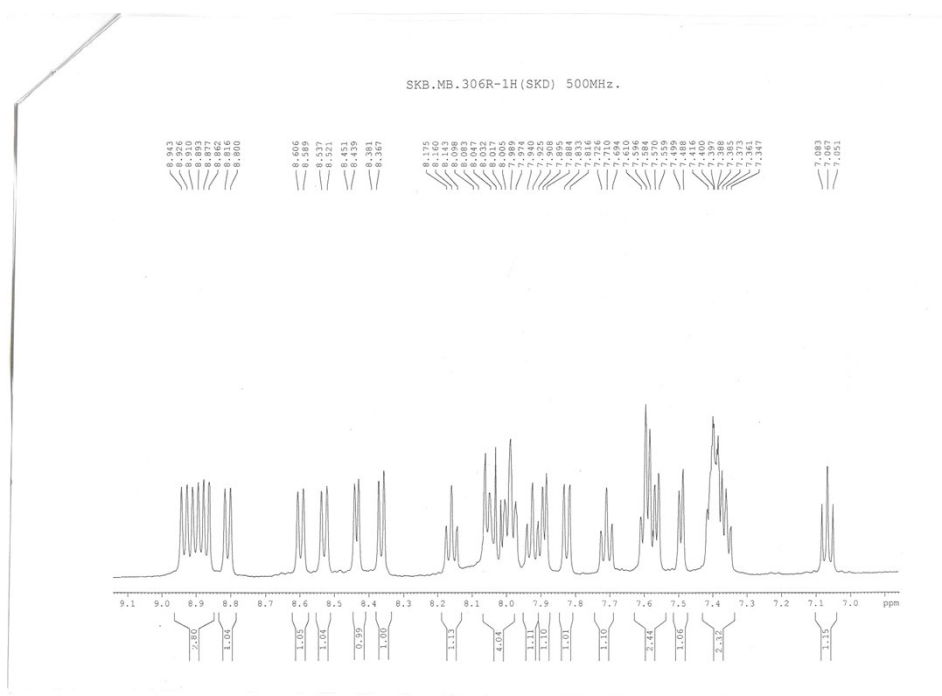
**Fig. S4**  $^1\text{H}$ - $^1\text{H}$  COSY NMR spectrum of **1** in  $\text{DMSO-}d_6$ .



**Fig. S5**  $^1\text{H}$ - $^1\text{H}$  COSY NMR spectrum of **2** in  $\text{DMSO}-d_6$ .

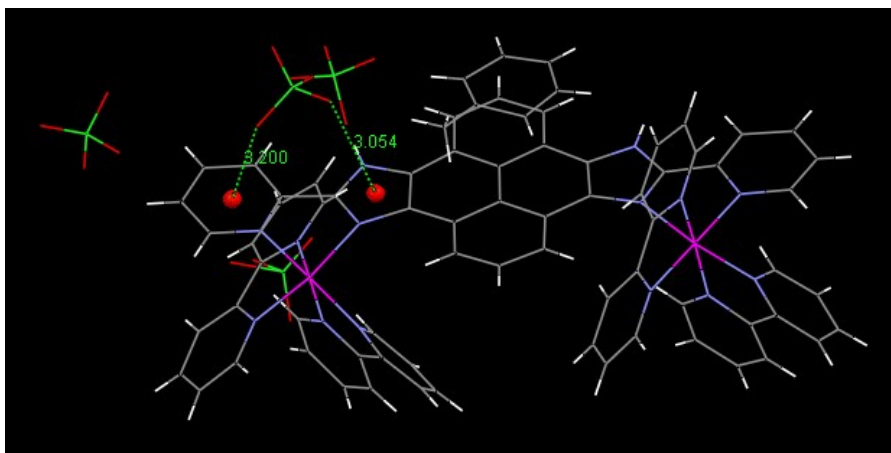


**Fig. S6** Scanned  $^1\text{H}$  NMR spectrum of complex **1** in  $\text{DMSO}-d_6$

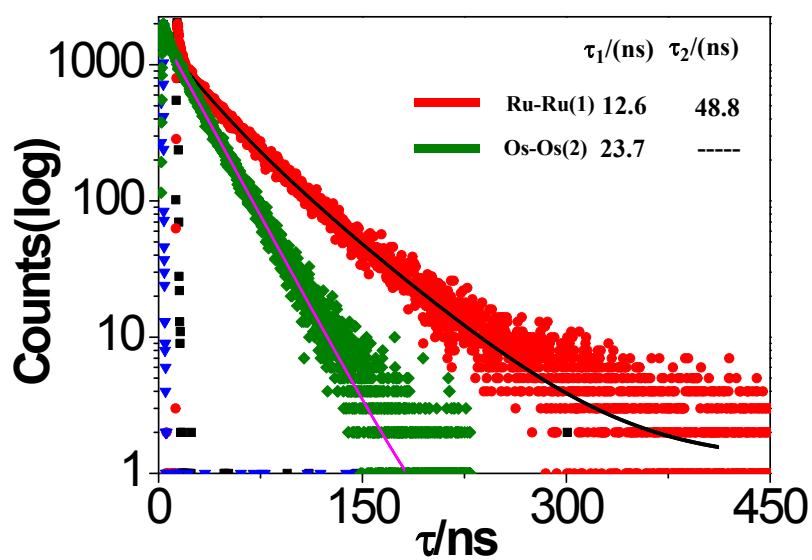


**Fig. S7** Scanned  $^1\text{H}$  NMR spectrum of complex **2** in  $\text{DMSO}-d_6$

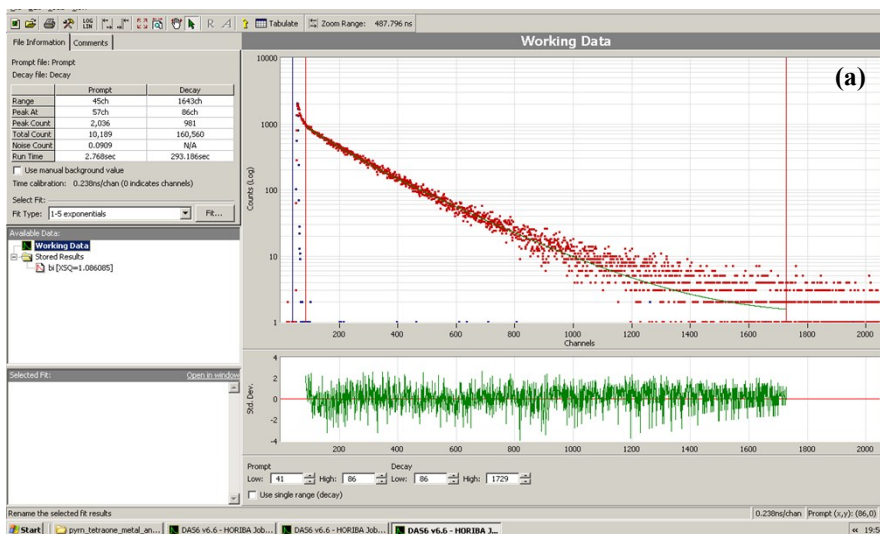




**Fig. S8** Occurrence of anion- $\pi$  interaction within **1**.



**Fig. S9** Time resolved decay profile of **1** and **2** in acetonitrile in their free state. Lifetimes of the complexes were acquired following excitation at 450 nm.



The fitted parameters are:

Hi reduced to: 1672 ch

SHIFT = 3.661401E-02 ch  
8.725042E-12 sec  
S.Dev = 1.132722E-10 sec

T1 = 53.05135 ch  
1.264203E-08 sec  
S.Dev = 1.437123E-09 sec

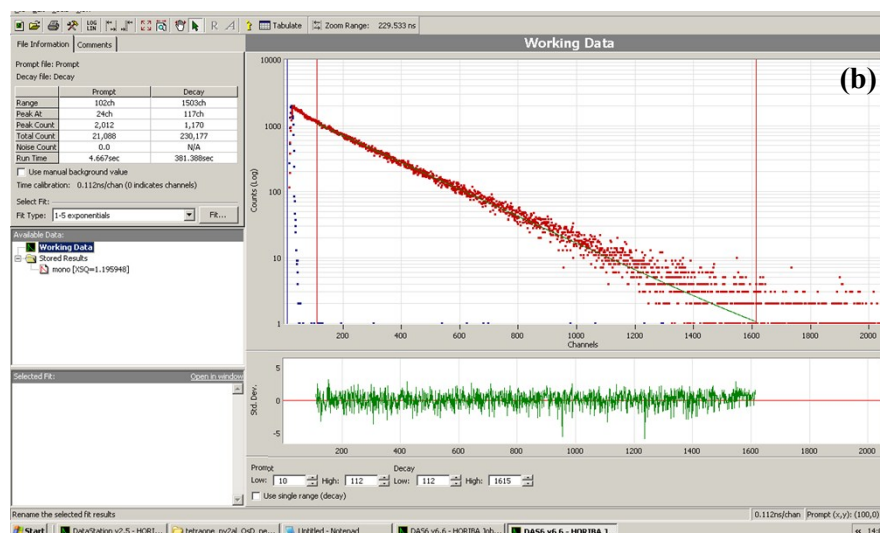
T2 = 199.9002 ch  
4.88358E-08 sec  
S.Dev = 2.417946E-10 sec

A = 1.617484  
S.Dev = 6.625398E-02

B1 = 4.368845E-02  
[12.51 Rel.Ampl]  
S.Dev = 1.003443E-03

B2 = 8.110892E-02  
[87.49 Rel.Ampl]  
S.Dev = 3.004865E-04

CHISQ = 1.144919  
[ 1595 degrees of freedom ]



The fitted parameters are:

SHIFT = 0.4 ch  
4.485248E-11 sec  
S.Dev = 4.485248E-11 sec

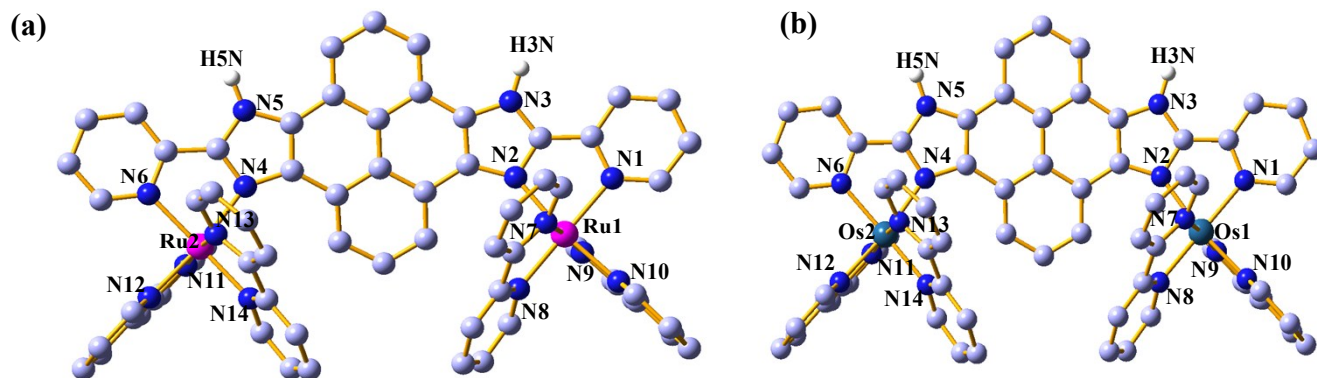
T1 = 212.1192 ch  
2.378518E-08 sec  
S.Dev = 6.840385E-11 sec

A = 0.1609985  
S.Dev = 0.0713059

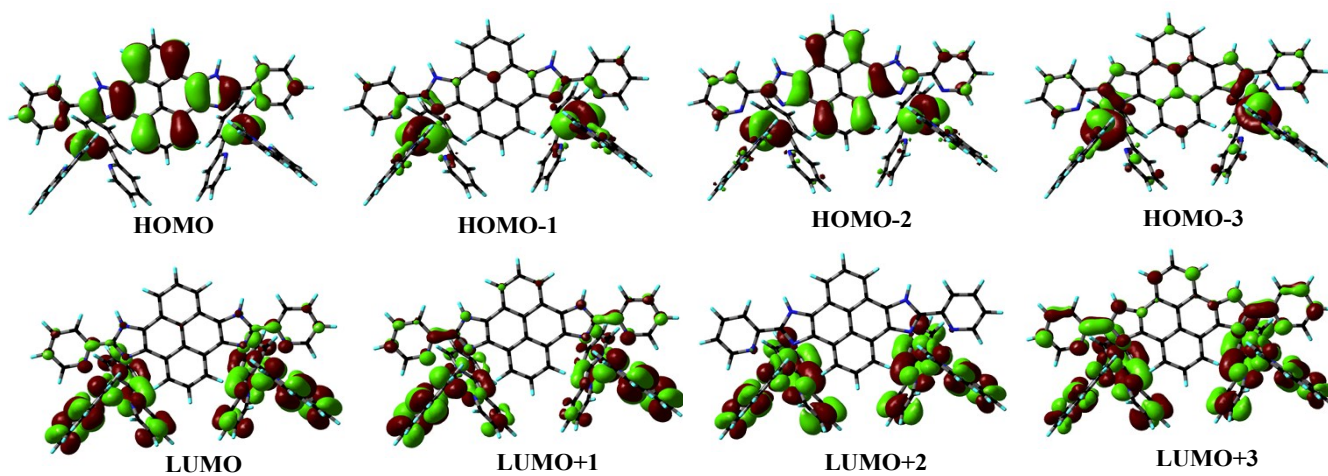
B1 = 7.701083E-02  
[100.00 Rel.Ampl]  
S.Dev = 1.650364E-04

CHISQ = 1.195948  
[ 1500 degrees of freedom ]

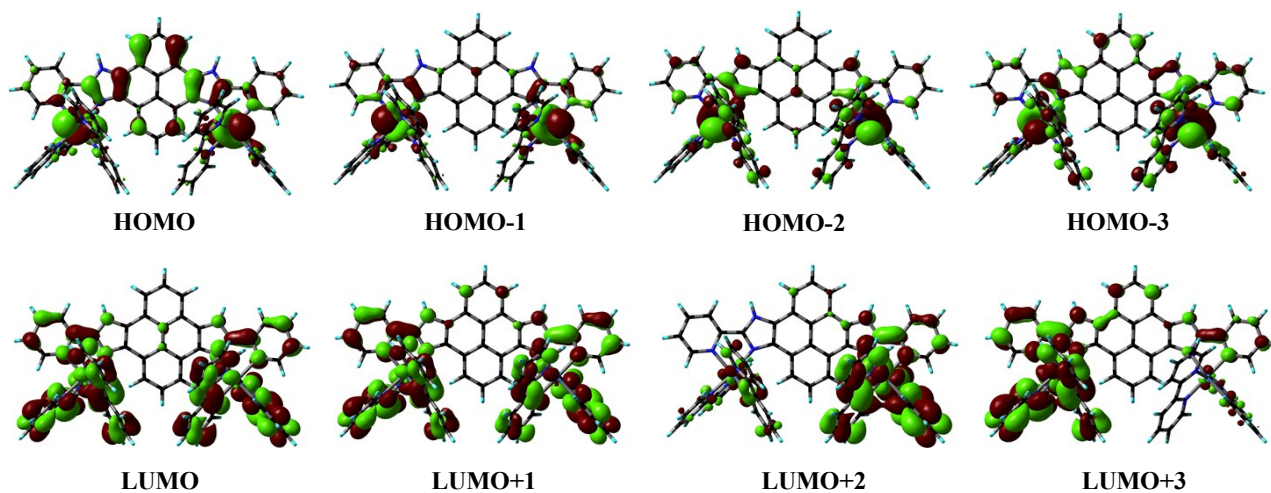
**Fig. S10** Fitted images and corresponding statistics of fit for the luminescence decay profile of **1** (a) and **2**(b) respectively in acetonitrile in their free states.



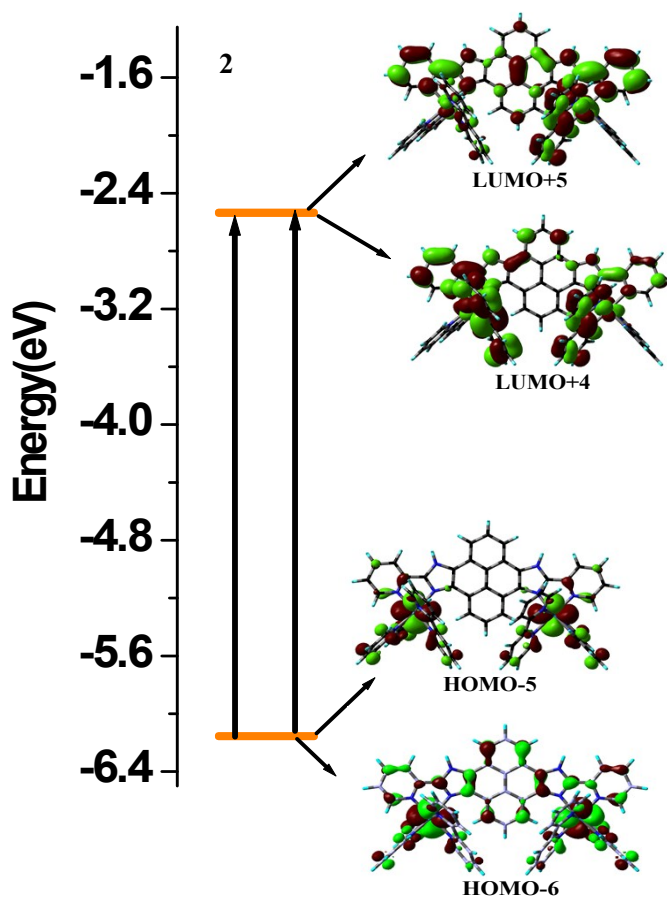
**Fig. S11** Optimized geometries and labelling scheme for (a)  $[(\text{bpy})_2\text{Ru}(\text{H}_2\text{Imz}_2\text{PPy}_2)\text{Ru}(\text{bpy})_2]^{4+}$  (**1**) and (b)  $[(\text{bpy})_2\text{Os}(\text{H}_2\text{Imz}_2\text{PPy}_2)\text{Os}(\text{bpy})_2]^{4+}$  (**2**) in solution phase.



**Fig. S12** Schematic drawings of the selective frontier molecular orbitals for  $[(\text{bpy})_2\text{Ru}(\text{H}_2\text{Imz}_2\text{PPy}_2)\text{Ru}(\text{bpy})_2]^{4+}$  (**1**) in solution phase.

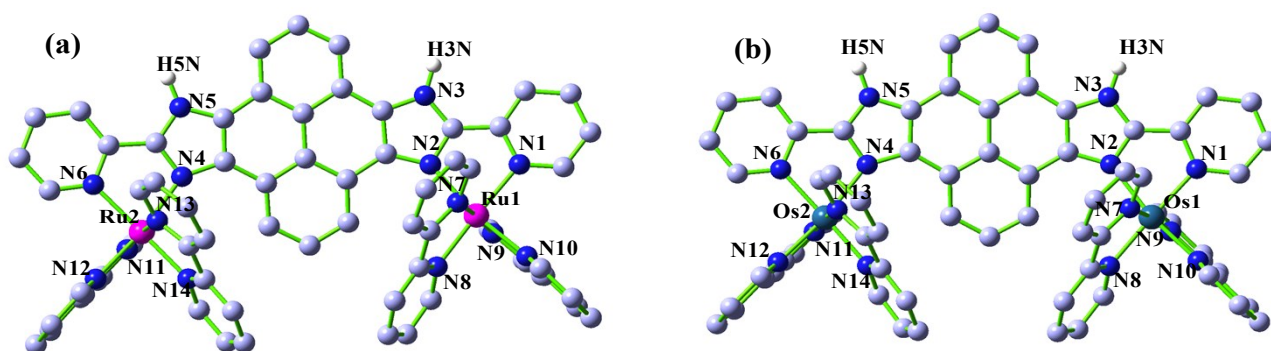


**Fig. S13** Schematic drawings of the selective frontier molecular orbitals for  $[(bpy)_2Os(H_2Imz_2PPy_2)Os(bpy)_2]^{4+}$  (**2**) in solution phase.

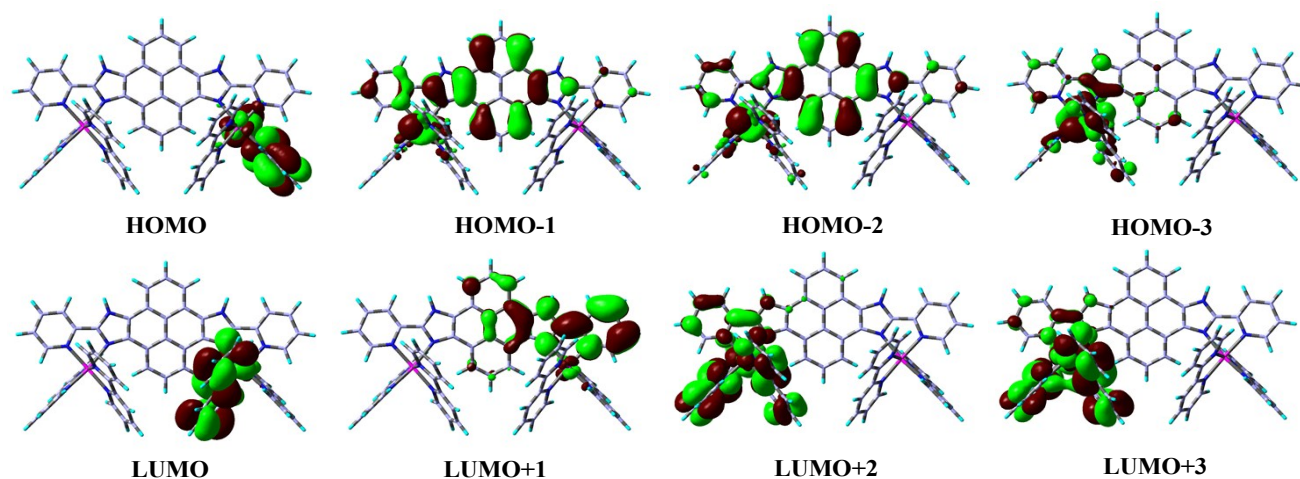


**Fig. S14** Energy level diagrams depicting the dominant transitions that comprise the lowest-energy absorption band for  $[(bpy)_2Os(H_2Imz_2PPy_2)Os(bpy)_2]^{4+}$  (**2**) in acetonitrile.

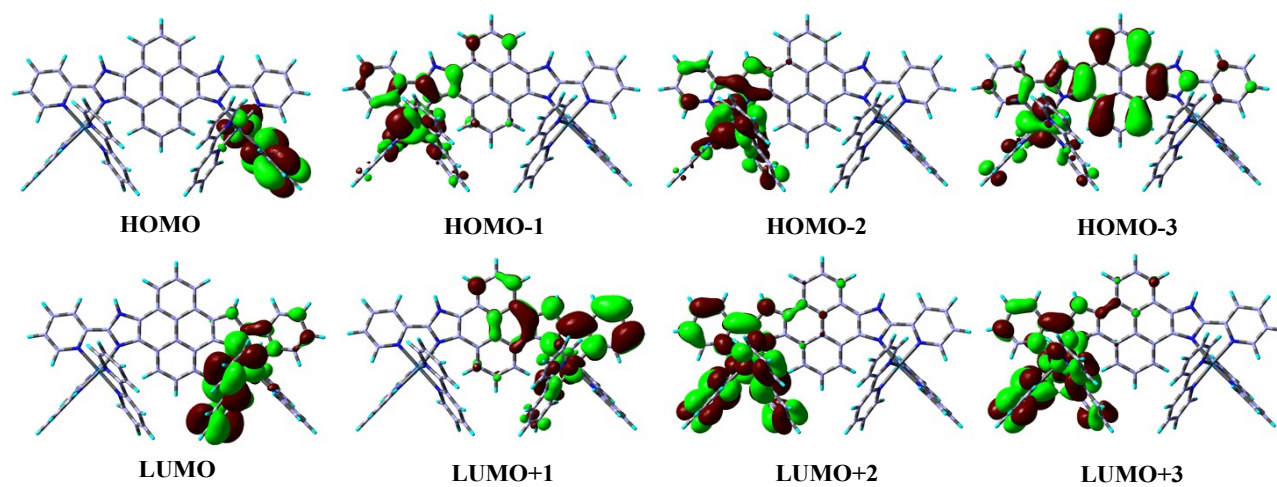




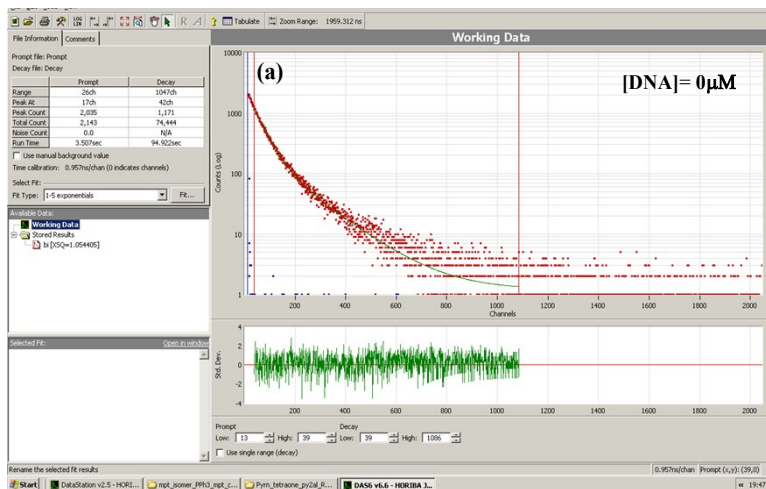
**Fig. S15** Optimized geometries and labelling scheme for  $[(\text{bpy})_2\text{Ru}(\text{H}_2\text{Imz}_2\text{PPy}_2)\text{Ru}(\text{bpy})_2]^{4+}$  (**1**) and  $[(\text{bpy})_2\text{Os}(\text{H}_2\text{Imz}_2\text{PPy}_2)\text{Os}(\text{bpy})_2]^{4+}$  (**2**) in UKS state.



**Fig. S16** Schematic drawings of the selective frontier molecular orbitals for  $[(\text{bpy})_2\text{Ru}(\text{H}_2\text{Imz}_2\text{PPy}_2)\text{Ru}(\text{bpy})_2]^{4+}$  (**1**) in UKS optimized state.



**Fig. S17** Schematic drawings of the selective frontier molecular orbitals for  $[(bpy)_2Os(H_2Imz_2PPy_2)Os(bpy)_2]^{4+}$  (**2**) in UKS optimized state.



The fitted parameters are:

SHIFT = 1.9 ch  
 1.818609E-09 sec  
 S.Dev = 1.69639E-09 sec

T1 = 42.97166 ch  
 4.143087E-08 sec  
 S.Dev = 7.184974E-10 sec

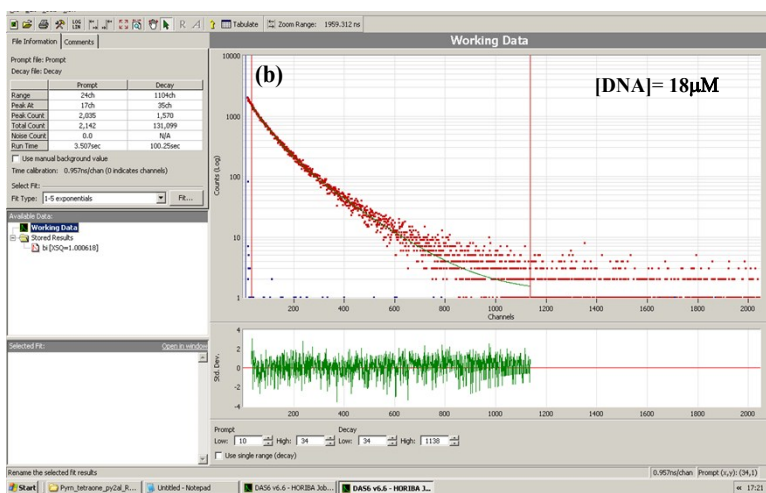
T2 = 144.4753 ch  
 1.341863E-07 sec  
 S.Dev = 1.83166E-09 sec

A = 1.023088  
 S.Dev = 5.134795E-02

B1 = 0.7422737  
 [66.34 Rel.Ampl]  
 S.Dev = 5.226232E-03

B2 = 0.1120418  
 [33.66 Rel.Ampl]  
 S.Dev = 1.130217E-03

CHISQ = 1.008743  
 [ 1253 degrees of freedom ]



The fitted parameters are:

SHIFT = -8.214164E-02 ch  
 -7.862292E-11 sec  
 S.Dev = 3.9433E-09 sec

T1 = 59.02127 ch  
 5.429296E-08 sec  
 S.Dev = 1.411408E-09 sec

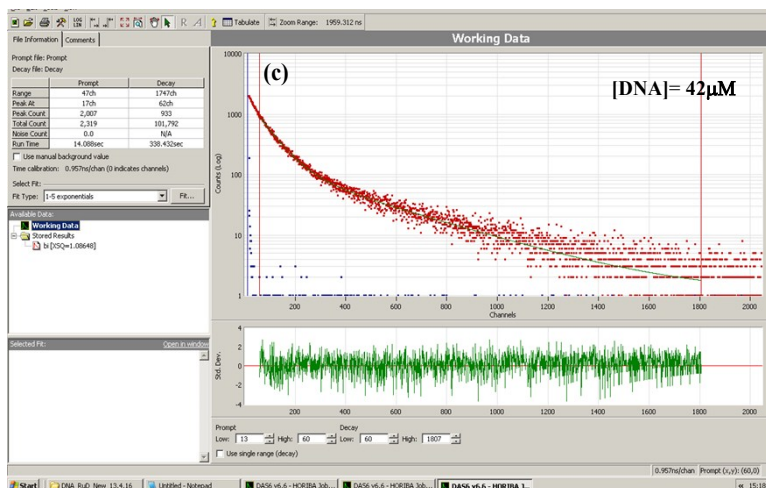
T2 = 151.5714 ch  
 1.650785E-07 sec  
 S.Dev = 1.572596E-09 sec

A = 1.230422  
 S.Dev = 7.701985E-02

B1 = 0.6455741  
 [51.68 Rel.Ampl]  
 S.Dev = 5.272183E-03

B2 = 0.2350785  
 [48.32 Rel.Ampl]  
 S.Dev = 1.822528E-03

CHISQ = 1.000618  
 [ 1099 degrees of freedom ]



The fitted parameters are:

SHIFT = 2.0327 ch  
 1.945625E-09 sec  
 S.Dev = 1.285637E-09 sec

T1 = 67.54663 ch  
 6.635312E-08 sec  
 S.Dev = 8.873542E-10 sec

T2 = 335.6617 ch  
 3.352829E-07 sec  
 S.Dev = 4.446409E-09 sec

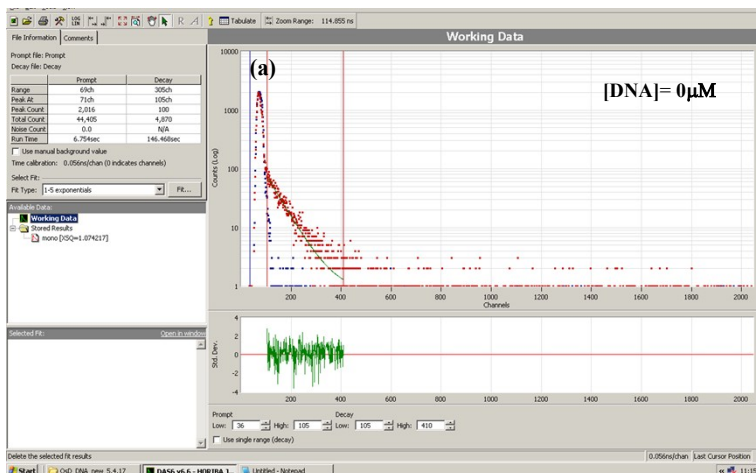
A = 1.026559  
 S.Dev = 6.958518E-02

B1 = 0.6158694  
 [65.10 Rel.Ampl]  
 S.Dev = 3.710504E-03

B2 = 6.645432E-02  
 [34.90 Rel.Ampl]  
 S.Dev = 5.109302E-04

CHISQ = 1.08648  
 [ 1742 degrees of freedom ]

**Fig. S18** Fitted images and corresponding statistics for the fit of each of the luminescence decay of **1** (a-c) in presence of increasing amount of CT-DNA in Tris-NaCl buffer medium (pH=7.30).



The fitted parameters are:

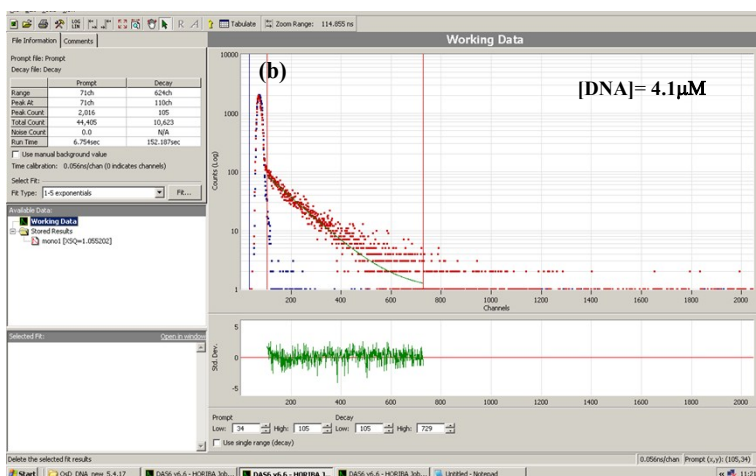
$$\begin{aligned} \text{SHIFT} &= -6.594157\text{E-}02 \text{ ch} \\ &= -3.699905\text{E-}12 \text{ sec} \\ \text{S.Dev} &= 1.261805\text{E-}10 \text{ sec} \end{aligned}$$

$$\begin{aligned} T1 &= 60.13293 \text{ ch} \\ &= 3.373989\text{E-}09 \text{ sec} \\ \text{S.Dev} &= 1.033712\text{E-}10 \text{ sec} \end{aligned}$$

$$\begin{aligned} A &= 0.8479583 \\ \text{S.Dev} &= 0.1325293 \end{aligned}$$

$$\begin{aligned} B1 &= 2.678487\text{E-}03 \\ &= [100.00 \text{ Rel.Amp}] \\ \text{S.Dev} &= 4.678271\text{E-}05 \end{aligned}$$

$$\begin{aligned} \text{CHISQ} &= 1.074217 \\ &= [ 302 \text{ degrees of freedom} ] \end{aligned}$$



The fitted parameters are:

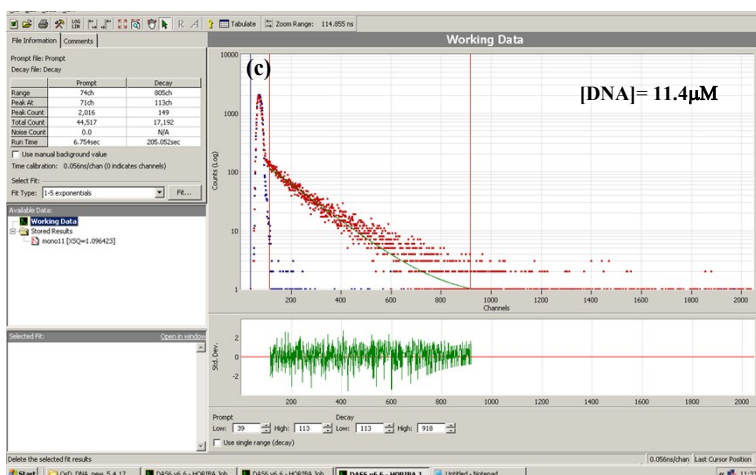
$$\begin{aligned} \text{SHIFT} &= -0.1310402 \text{ ch} \\ &= -7.352511\text{E-}12 \text{ sec} \\ \text{S.Dev} &= 3.349409\text{E-}10 \text{ sec} \end{aligned}$$

$$\begin{aligned} T1 &= 111.1268 \text{ ch} \\ &= 6.235195\text{E-}09 \text{ sec} \\ \text{S.Dev} &= 1.135144\text{E-}10 \text{ sec} \end{aligned}$$

$$\begin{aligned} A &= 0.9556906 \\ \text{S.Dev} &= 8.709306\text{E-}02 \end{aligned}$$

$$\begin{aligned} B1 &= 2.517672\text{E-}03 \\ &= [100.00 \text{ Rel.Amp}] \\ \text{S.Dev} &= 2.888786\text{E-}05 \end{aligned}$$

$$\begin{aligned} \text{CHISQ} &= 1.055202 \\ &= [ 621 \text{ degrees of freedom} ] \end{aligned}$$



The fitted parameters are:

$$\begin{aligned} \text{SHIFT} &= -5.846859\text{E-}02 \text{ ch} \\ &= -3.280605\text{E-}12 \text{ sec} \\ \text{S.Dev} &= 2.832513\text{E-}10 \text{ sec} \end{aligned}$$

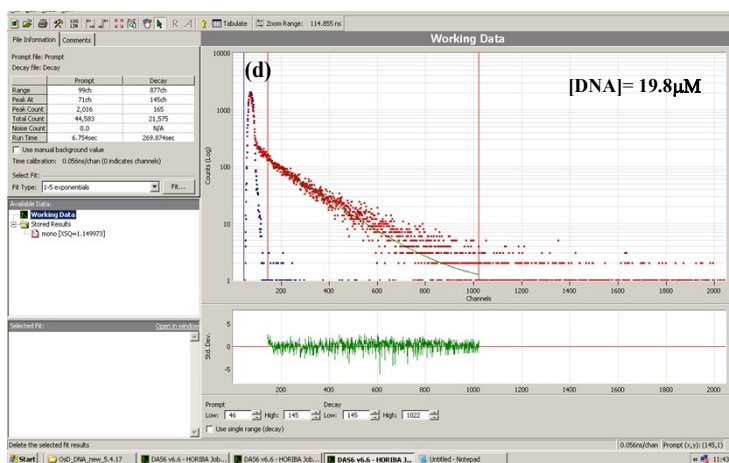
$$\begin{aligned} T1 &= 135.5538 \text{ ch} \\ &= 7.605765\text{E-}09 \text{ sec} \\ \text{S.Dev} &= 1.075715\text{E-}10 \text{ sec} \end{aligned}$$

$$\begin{aligned} A &= 0.7150995 \\ \text{S.Dev} &= 7.388485\text{E-}02 \end{aligned}$$

$$\begin{aligned} B1 &= 3.48344\text{E-}03 \\ &= [100.00 \text{ Rel.Amp}] \\ \text{S.Dev} &= 3.001725\text{E-}05 \end{aligned}$$

$$\begin{aligned} \text{CHISQ} &= 1.096423 \\ &= [ 802 \text{ degrees of freedom} ] \end{aligned}$$





The fitted parameters are:

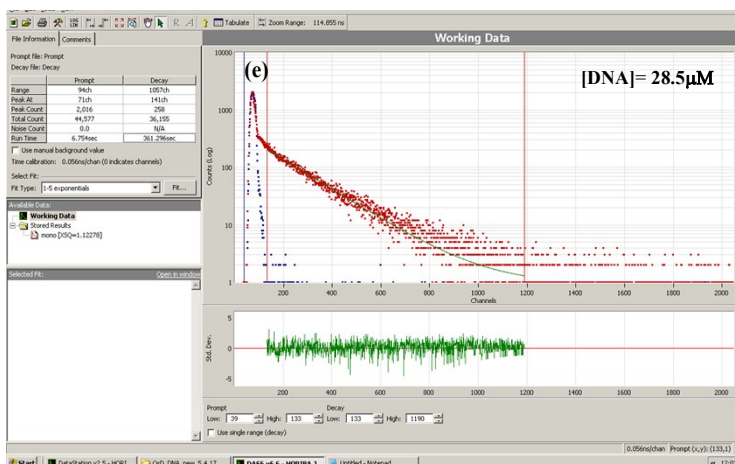
SHIFT = -9.999993E-02 ch  
-5.61088E-12 sec  
S.Dev = 8.336415E-10 sec

T1 = 153.2918 ch  
8.601025E-09 sec  
S.Dev = 1.121407E-10 sec

A = 0.8411134  
S.Dev = 7.882145E-02

B1 = 4.62003E-03  
[100.00 Rel.Ampl]  
S.Dev = 3.580114E-05

CHISQ = 1.149973  
[ 874 degrees of freedom ]



The fitted parameters are:

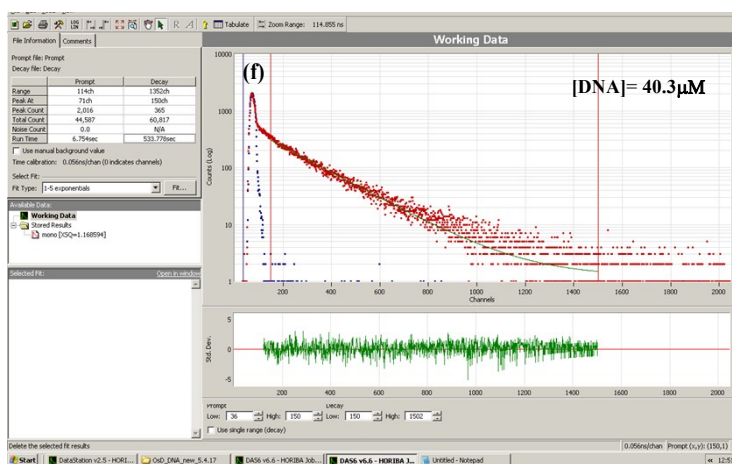
SHIFT = -9.999999E-02 ch  
-5.610883E-12 sec  
S.Dev = 7.500983E-10 sec

T1 = 165.403 ch  
9.280567E-09 sec  
S.Dev = 8.477425E-11 sec

A = 0.953526  
S.Dev = 7.079282E-02

B1 = 6.580617E-03  
[100.00 Rel.Ampl]  
S.Dev = 3.789648E-05

CHISQ = 1.12278  
[ 1054 degrees of freedom ]



The fitted parameters are:

SHIFT = -0.1 ch  
-5.610884E-12 sec  
S.Dev = 6.517638E-10 sec

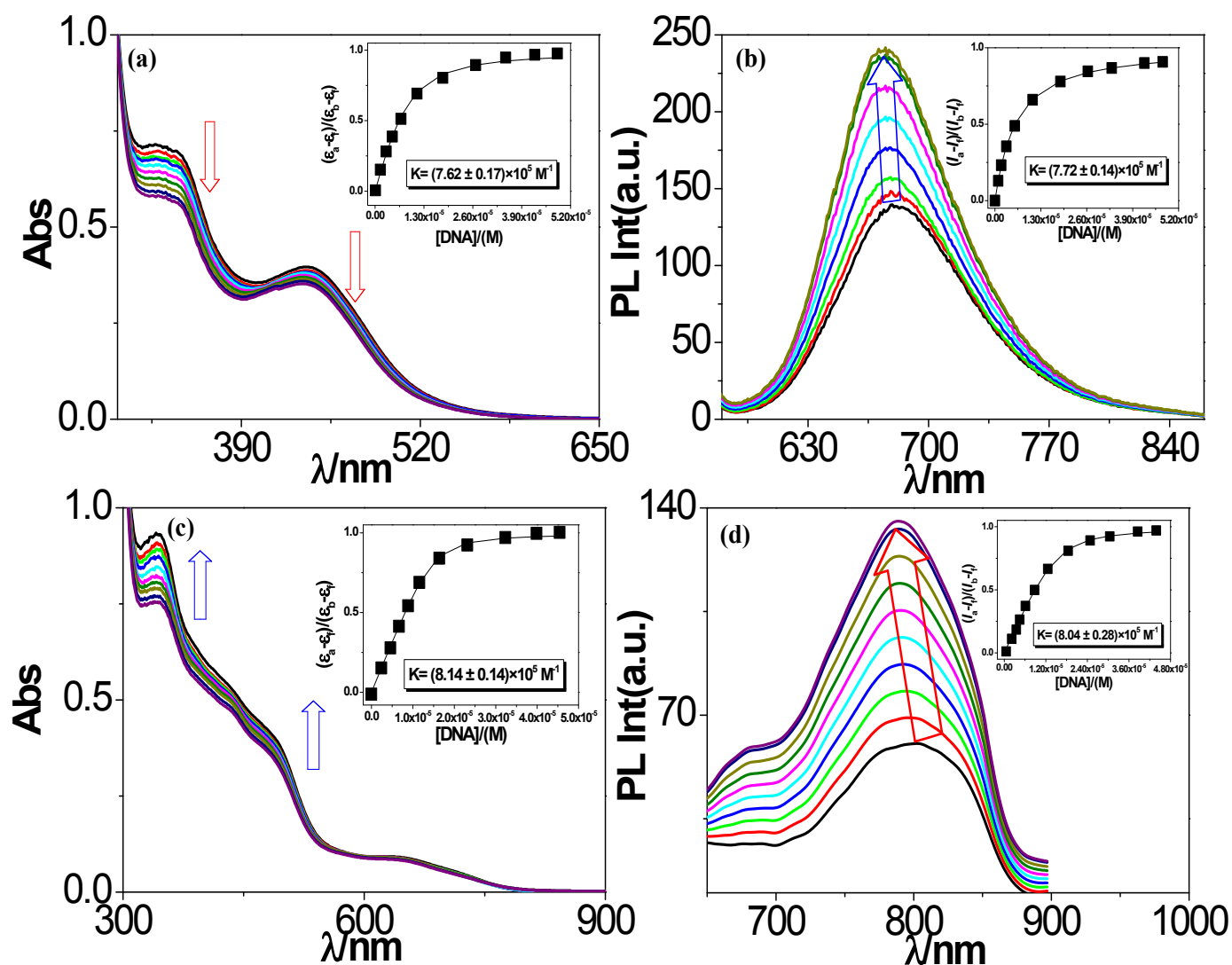
T1 = 190.7604 ch  
1.070335E-08 sec  
S.Dev = 6.548448E-11 sec

A = 1.167494  
S.Dev = 5.536998E-02

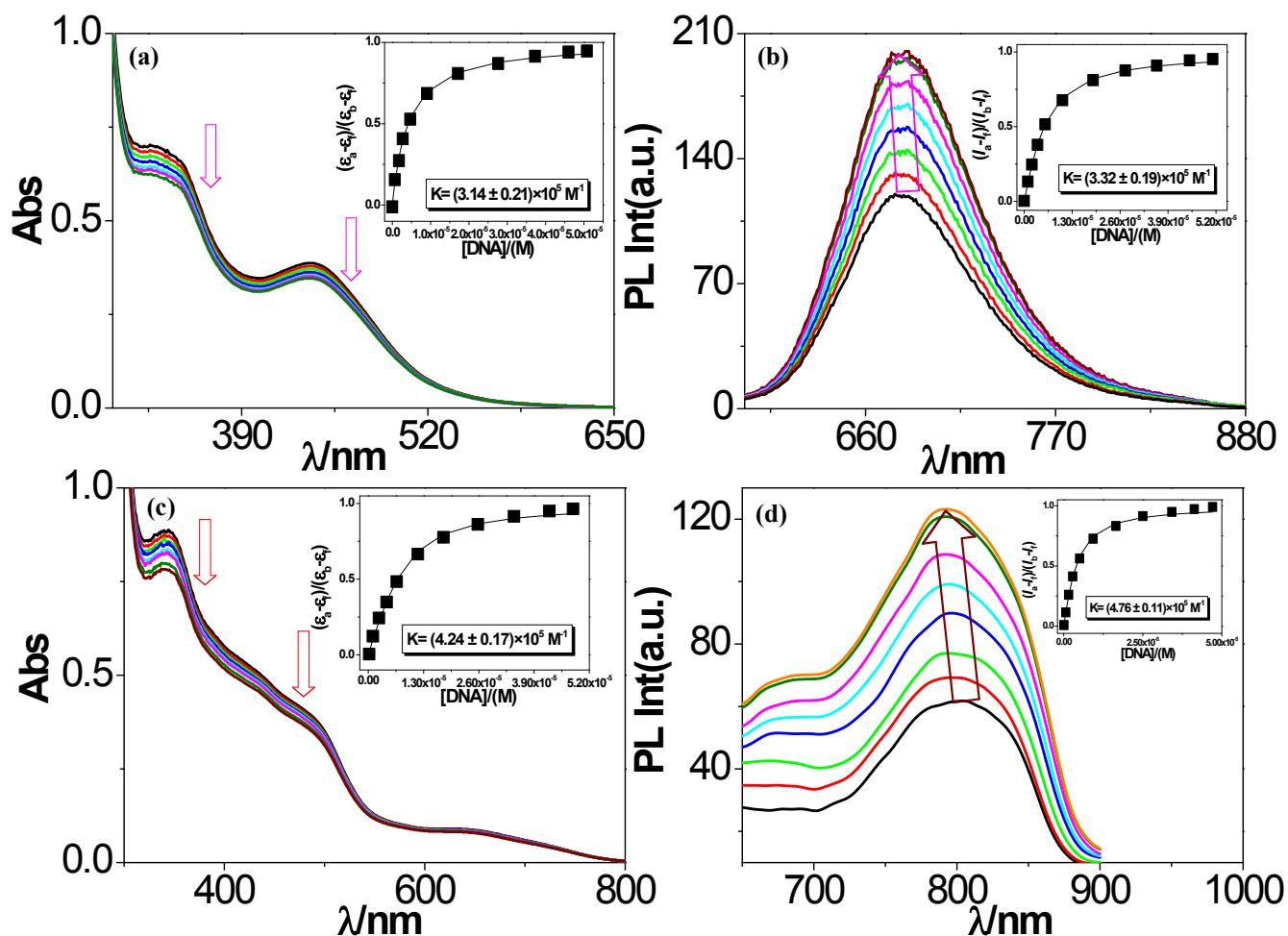
B1 = 1.007874E-02  
[100.00 Rel.Ampl]  
S.Dev = 4.354247E-05

CHISQ = 1.168594  
[ 1453 degrees of freedom ]

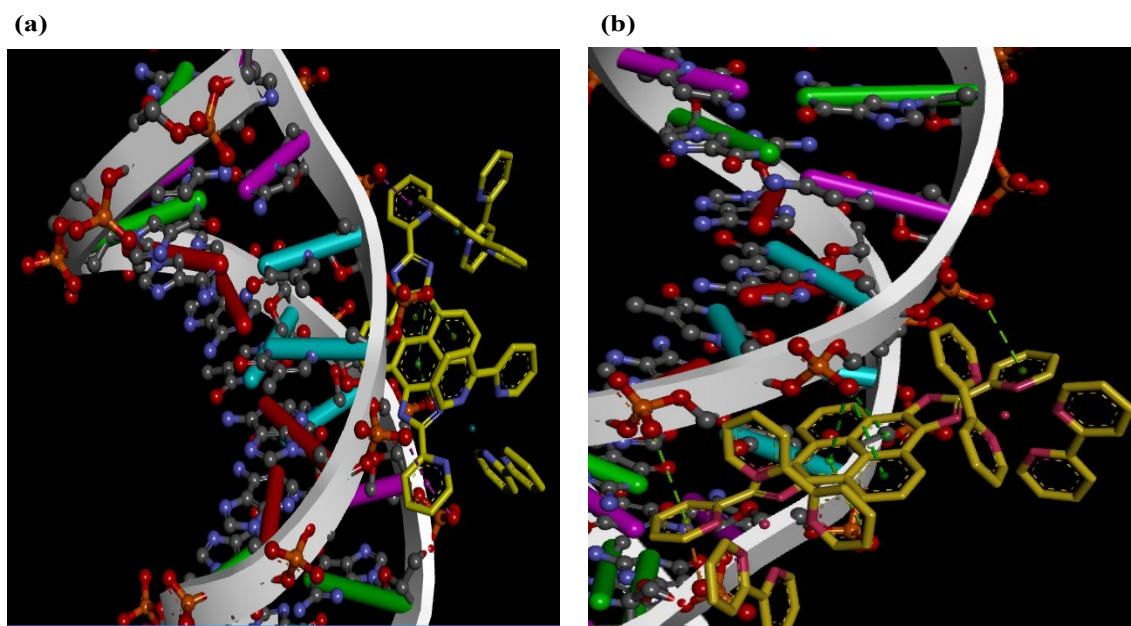
**Fig. S19** Fitted images and corresponding statistics for the fit of each of the luminescence decay of **2** (a-f) in presence of increasing amount of CT-DNA in Tris-NaCl buffer medium (pH=7.30).



**Fig. S20** Changes in UV-vis absorption (a, c) and luminescence spectra (b, d) of **1** and **2** (20  $\mu$ M) in presence of increasing amount of CT-DNA (0–40  $\mu$ M for **1** and 0–36  $\mu$ M for **2**) in 300 mM NaCl-Tris buffer medium (pH=7.30). Insets show the binding profile with DNA. Excitation wavelength used for acquiring the luminescence spectra is 450 nm for **1** and 480 nm for **2**.



**Fig. S21** Changes in UV-vis absorption (a, c) and luminescence spectra (b, d) of **1** and **2** (20 μM) in presence of increasing amount of CT-DNA (0-45 μM for **1** and 0-40 μM for **2**) in 1M NaCl-Tris buffer medium (pH=7.30). Insets show the binding profile with DNA. Excitation wavelength used for acquiring the luminescence spectra is 450 nm for **1** and 480 nm for **2**.



**Fig. S22** Occurrence of anion- $\pi$  interaction between DNA and the complexes; (a) for **1** and (b) for **2**.

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