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Electronic Supplementary Information

Cycloplatinated(II) complexes bearing O,S-heterocyclic ligand: searching for anticancer drugs

Masood Fereidoonnezhad,^a Zahra Ramezani,^a Mahshid Nikravesh,^b Jalalaldin Zangeneh,^a Mohsen Golbon Haghighi,^c Zahra Faghih,^d Behrouz Notash,^c and Hamid R. Shahsavari^b*

^aCancer, Environmental and Petroleum Pollutants Research Center; Department of Medicinal Chemistry, School of Pharmacy, Ahvaz Jundishapur University of Medical Sciences, Ahvaz, Iran.

^bDepartment of Chemistry, Institute for Advanced Studies in Basic Sciences (IASBS), Zanjan 45137-66731, Iran.

^cDepartment of Chemistry, Shahid Beheshti University, Evin, Tehran 19839-69411, Iran.

^dShiraz Institute for Cancer Research, School of Medicine, Shiraz University of Medical Sciences, Shiraz, Iran.

Email: shahsavari@iasbs.ac.ir

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Figure S1. ¹H NMR (down) and HH COSY (up) spectra of **2a** in CD₂Cl₂ at room temperature. The signals assignments are depicted and the platinum satellites are indicated by *.





Figure S3. DEPT 135° spectrum of 2a in CD₂Cl₂.



Figure S4. 195 Pt{ 1 H} NMR spectrum of 2a in CD₂Cl₂.







Figure S6. ¹³C{¹H} NMR spectrum of **2b** in CD₂Cl₂.



Figure S7. DEPT 135° spectrum of 2b in CD₂Cl₂.



Figure S8. ¹⁹⁵Pt{¹H} NMR spectrum of 2b in CD₂Cl₂.



Figure S9. The calculated possible structures for SpyO ligand while it coordinated to 1.

Table S1. Crystanographic	c and structure refinement data for 2a.
Formula	C16H12N2OPtS
Formula weight	475.42
T (K)	298(2)
λ (Å)	0.71073
Crystal system	Monoclinic
Space group	$P2_{1/n}$
Crystal size(mm)	0.20 imes 0.10 imes 0.10
<i>a</i> (Å)	8.6392(17)
<i>b</i> (Å)	8.6163(17)
<i>c</i> (Å)	20.066(4)
α (°)	90
β (°)	93.80(3)
γ (°)_	90
$V(\text{\AA}^3)$	1490.4(5)
Ζ	4
$D_{\text{calc}} (\text{g cm}^{-1})$	2.119
hetamin, $ heta$ max (°)	2.57 - 25.00
F000	896
$\mu (\text{mm}^{-1})$	9.553
Index ranges	$-8 \le h \le 10$
	$-10 \le k \le 10$
	$-23 \le l \le 23$
Data collected	7906
Unique data	2624
$R_{I}^{a}, wR_{2}^{D} (I > 2\sigma (I))$	0.0567, 0.0728
R_1^a , wR_2^b (all data)	0.1168, 0.0835
GOF on F^2 (S)	0.947
CCDC No.	1568889
$a_{\mathbf{D}}$ SHELLEHASE b	D $(\Sigma (-\Sigma 5 - \Sigma 2)^2) / \Sigma (\Sigma 2)^2 1 / 2$

 Table S1. Crystallographic and structure refinement data for 2a.

 ${}^{a}R_{1} = \Sigma ||F_{o}| - |F_{c}|| / \Sigma |F_{o}|, {}^{b}wR_{2} = [\Sigma (w(F_{o}{}^{5} - F_{c}{}^{2})^{2}) / \Sigma w(F_{o}{}^{2})^{2}]^{1/2}$



Figure S10. The crystal packing of 2a.



Figure S11. Complex **2a** is displaying the intermolecular contacts. The supramolecular packing is formed by dimers supported by intermolecular $\pi \cdots \pi$ interactions involving two SpyO ligands.

Energies(eV)	Number	Pt	SpyO	рру
-0.798	LUMO+3	1	98	1
-1.017	LUMO+2	2	2	97
-1.365	LUMO+1	1	97	2
-1.672	LUMO	6	2	91
-5.487	НОМО	36	45	19
-5.871	HOMO-1	47	28	25
-6.286	HOMO-2	94	2	3
-6.470	НОМО-3	8	8	84

Table S2. Fragment contributions (%; from atomic orbital contributions) to the frontier orbitals of **2a** in CH₂Cl₂ solution.

Table S3. Fragment contributions (%; from atomic orbital contributions) to the frontier orbitals of **2b** in CH₂Cl₂ solution.

Energies(eV)	Number	Pt	SpyO	bzq
-0.810	LUMO+3	1	98	1
-1.216	LUMO+2	3	11	86
-1.381	LUMO+1	1	91	8
-1.844	LUMO	4	2	94
-5.433	НОМО	33	37	30
-5.784	HOMO-1	44	35	22
-6.296	HOMO-2	94	3	4
-6.425	HOMO-3	10	17	73

state	Monoexcitations ^a		λ_{cal}/nm	oscillator	main character
		E/eV		strength	
S 1	H→L (63%)	3.190	389	0.114	MLCT/LC/L'LCT
S2	H-1→L (60%)	3.364	369	0.078	MLCT/LC/L'LCT
	H→L (31%)				
	H→L+1 (17%)				
S 8	H-1→L+2 (56%)	4.043	307	0.067	MLCT/LC/L'LCT
	H→L+3 (22%)				
S10	H→L+3 (58%)	4.111	302	0.077	MLCT/LC/L'LCT
	H-3→L (20%)				
S14	H-1→L+3 (66%)	4.404	282	0.105	MLCT/LC/L'LCT
	H-2→L+4 (11%)				

Table S4. Selected vertical singlet excitations of **2a** from TD-DFT calculations at the ground state geometry in CH₂Cl₂ solution (M = Pt, L = ppy, L' = SpyO).

Table S5. Selected vertical singlet excitations of **2b** from TD-DFT calculations at the ground state geometry in CH₂Cl₂ solution (M = Pt, L = bzq, L' = SpyO).

state	Monoexcitations ^a		λ_{cal}/nm	oscillator	main character
		E/eV		strength	
S 1	H→L (66%)	3.019	411	0.094	MLCT/LC/L'LCT
	H-1→L (21%)				
S 3	H→L+1 (67%)	3.479	356	0.065	MLCT/LC/L'LCT
	H-1→L (16%)				
S5	H→L+2 (67%)	3.618	343	0.146	MLCT/LC/L'LCT
	H-4→L (12%)				
	H-1→L+1 (11%)				
	H-1→L (10%)				
S 9	H-3→L (50%)	4.006	310	0.098	LC
	H-2→L+4 (11%)				
S11	H→L+3 (56%)	4.081	304	0.084	MLCT/LC/L'LCT
	H-1→L+1 (24%)				
	H-3→L (17%)				
S13	H-4→L (59%)	4.181	297	0.073	LC/L'LCT
	H-3→L+2 (28%)				
	H-3→L+1 (15%)				





Figure S14. 3D ligand-receptor interactions of 2a with DNA (PDB code: 1BNA).



Figure S15. 3D ligand-receptor interactions of 2b with DNA (PDB code: 1BNA).



Figure S16. 3D ligand-receptor interactions of 2a with DNA (PDB code: 198D).



Figure S17. 3D ligand-receptor interactions of 2b with DNA (PDB code: 198D).



Figure S18. 3D ligand-receptor interactions of 2a with DNA (PDB code: 1LU5).



Figure S19. 3D ligand-receptor interactions of 2b with DNA (PDB code: 1LU5).