Palladium(II) induced complete conformational enrichment of the syn isomer of N,N'-Bis(4-pyridylformyl)piperazine

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Contents:

The file includes ¹H, ¹³C NMR, H-H COSY, C-H COSY of L and 1 (in CDCl₃ and D₂O for L; and in D₂O for 1). ¹H NMR spectrum of 1' in D₂O. Variable temperature ¹H NMR of L in D₂O. Stacking diagram of ¹H NMR of L in CDCl₃, D₂O and DMSO-d₆. ESI MS for complex 1', crystallographic information table for 1' and L, ORTEP for 1'and L, energy minimized structures for *syn* and *anti* forms of L, complexes 1 and 2, powder XRD data for pre-crystallized and crystallized L and the comparison with simulated spectrum.



Fig. S1 400 MHz 1 H NMR for **L** in CDCl₃.



Fig. S2 ¹³C NMR for L in CDCl₃.



Fig. S3 H-H COSY for \mathbf{L} in CDCl₃



Fig. S4 C-H COSY for L in $CDCl_3$



Fig. S5 400 MHz 1 H NMR for **L** in D₂O.



Fig. S6 13 C NMR for **L** in D₂O.



Fig. S7 H-H COSY for \mathbf{L} in D_2O .



Fig. S8 Expansion of H-H COSY for L in D_2O .



Fig. S9 C-H COSY for \mathbf{L} in D_2O .



Fig. S10 Expansion of C-H COSY for L in D₂O.



Fig. S11 400 MHz 1 H NMR for **1** in D₂O.



Fig. S12 13 C NMR for **1** in D₂O.



Fig. S13 H-H COSY for 1 in D_2O .



Fig. S14 C-H COSY for 1 in D₂O.

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Fig. S15 400 MHz 1 H NMR for **1'** in D₂O.



Fig. S16 400 MHz ¹H NMR in D₂O for **L** at different temperatures; (i) 25 °C, (ii) 40 °C, (iii) 50 °C, (iv) 60 °C, (v) 70 °C, and (vi) 80 °C respectively.



Fig. S17 400 MHz ¹H NMR for **L** recorded in different solvents; in (i) CDCl₃, (ii) D₂O, and (iii) DMSO-d₆. The signals shown in red color belongs to the piperazine protons.



Fig. S18 ESI MS spectrum for complex **1**'showing isotopic distribution pattern for the peak at m/z=341 corresponding to $[1'-3ClO_4]^{3+}$.

Empirical formula	C36 H48 Cl4 N12 O23 Pd2
Formula weight	1371.46
Temperature (T)	293(2) K
Wavelength (λ)	0.71073 Å
Crystal system, space group	Orthorhombic, Cmc2(1)
Unit cell dimensions	
a = 18.8926(6) Å	
b = 17.6085(6) Å	
c = 17.0599(6) Å	
Volume (V)	5675.3(3) Å ³
Z, Calculated density	4, 1.605 mg/m^3
Absorption coefficient	0.906 mm ⁻¹
F(000)	2768
Crystal size	0.35 x 0.28 x 0.26 mm
Theta range for data 1	.58 to 25.00°.
collection	
Reflections collected/unique	32532 / 5142 [R(int) = 0.0262]
Completeness to theta $= 25.0$	0 100.0 %
Absorption correction	None
Max. and min. transmission	0.7985 and 0.7421
Refinement method	Full-matrix least-squares on F ²
Data / restraints / parameters	5142 / 5 / 369
Goodness-of-fit on F ² 1.122	
Final R indices [I>2sigma(I)]	R1 = 0.0405, wR2 = 0.1244
R indices (all data)	R1 = 0.0465, wR2 = 0.1341
Largest diff. peak and hole0.82	21 and -0.444 e.Å ⁻³

Table S1 Summary of X-ray crystallographic data collection and refinement parameters for 1'



Fig. S19 ORTEP diagram for complex **1**'. Hydrogen atoms, solvent molecules and anions are excluded for clarity. Thermal ellipsoids are shown in 50% probability level.

Table S2: Summary of X-ray crystallographic data collection and refinement parameters for L		
Empirical formula C16H16N4O2		
Formula weight 2	296.33	
Temperature 298(2) K		
Wavelength 0.71073 A		
Crystal system, space group Monoclinic, P2(1)/n		
Unit cell dimensions		
a = 9.1945(6) Å		
b = 8.2633(5) Å		
c = 9.9539(6) Å		
$\beta = 107.528(2)^{\circ}.$		
Volume (V)721.15(8) Å ³		
Z, Calculated density $4, 1.387 \text{ mg/m}^3$		
Absorption coefficient 0.125 mm ⁻¹		
F(000) 308		
Crystal size 0.25 x 0.20 x 0.15 mm		
Theta range for data collection 2.65 to 25.00 deg.		
Reflections collected / unique $4185 / 1211 [R_{(int)} = 0.0213]$		
Completeness to theta = 25.00 95.7 9	%	
Absorption correction Multi-scan		
Max. and min. transmission 0.98	315 and 0.9695	
Refinement method Full-matrix least-squares on F ²		
Data / restraints / parameters 1211 /	/ 0 / 101	
Goodness-of-fit on F^2 1.0	168	
Final R indices $[I > 2 \text{sigma}(I)]RI = 0.0367, wR2 = 0.1259$	9	
R indices (all data) $R1 = 0.0466, wR2 = 0.1378$		
Extinction coefficient 0.0	094(16)	
Largest diff. peak and hole0.184 and -0.148 e.Å ⁻³		



Fig. S20 ORTEP diagram for **L**. Hydrogen atoms are excluded for clarity. Thermal ellipsoids are shown in 50% probability level.



Fig. S21 Powder-XRD data for the ligand L; a) simulated from the single crystal X-ray data, (bc) experimental diffraction pattern for L b) after re-crystallization from water and c) after recrystallization from DCM



Fig. S22 Energy minimized structure for both *syn* and *anti*-conformations obtained by DFT calculations.



Fig. S23 Energy minimized structure for complexes of Pd(II) with L; a) in *syn*-conformation and b) in *anti*-conformation obtained by DFT calculations using B3LYP-LANL2DZ basis sets as included in *Gaussian 09* package¹ (corresponding thermodynamic parameters are listed below the structures).

Reference

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