

Supporting Information

A neutral Pt₃ stack unsupported by any bridging ligand

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Preparation of **1** and **2**.

Pt(Hpyzpy)Cl₂·H₂O (1a) and **Pt(Hpyzpy)Cl₂·(CH₃)₂CO (1b)**. K₂PtCl₄ (1.26 mmol, 521 mg) and Hpyzpy (0.627 mmol, 91 mg) were dissolved in water (150 mL) and the pH was brought to 3.7 by adding a few drops of 2 M HCl. Within 10 h at room temperature, two fractions (15 and 20 mg) of a yellow solid were filtered off, which contained a mixture of **1a** and *trans*-Pt(pyzyzpy)₂ (**2**), according to ¹H NMR spectra (Me₂SO-d₆). The third fraction (142 mg, 53 %) recovered by filtration after 6 d at RT proved to contain exclusively **1a**. Anal. Calcd (%) for C₈H₉N₃OCl₂Pt: C, 22.4; H, 2.1; N, 9.8. Found: C, 22.4; H, 2.2; N, 9.5. ¹H NMR (δ, ppm, CDCl₃): 11.7 (NH, b), 9.60 (dd, 5.6, 1.3 Hz; ¹⁹⁵Pt satellites ca 42 Hz; H6), 8.10 (ddd, 7.4, 7.4, 1.3 Hz; H4), 7.02 (d, 2.7 Hz; H5'), 7.77 (dd, 7.4, 1.3 Hz, H3), 7.51 (ddd, 7.4, 5.6, 1.3 Hz; H5), 6.92 (d, 2.7 Hz; H4'). Crude **1a** was dissolved in a minimum amount of DMF and acetone was added. Pale yellow crystals of Pt(Hpyzpy)Cl₂·(CH₃)₂CO (**1b**) were isolated in low yield and characterized by X-ray analysis.

trans-Pt(pyzyzpy)₂ (**2**). **1** (0.46 mmol, 200 mg) was suspended in water (400 mL), Hpyzpy (1.38 mmol, 200 mg) was added, and the mixture was stirred at 100 °C for 3 d. Yellow **2** was filtered off, washed with water and dried at 40 °C. The yield was 71 % based on **1**. Alternatively, **2** was obtained by refluxing an aqueous mixture of K₂PtCl₄ and Hpyzpy (1:3) for 48 h. Filtration of the yellow precipitate, washing with water and drying in air gave **2** in 77 % yield. Anal. Calcd (%) for C₁₆H₁₂N₆Pt: C, 39.8; H, 2.5; N, 17.4. Found: C, 39.6; H, 2.7; N, 17.6. Orange-yellow single crystals suitable for X-ray analysis were obtained upon recrystallization of the crude product from CHCl₃.

X-Ray crystallography

Intensity data collections of **1** and **2** were performed on an Enraf–Nonius Kappa CCD (compound **1**) and on an Oxford Diffraction Xcalibur (compound **2**) diffractometers. Data reduction and cell refinement were carried out using the DENZO and SCALE-PACK (for **1**) programs,¹ and CrysAlisPro software (for **2**).² All the structures were solved by direct

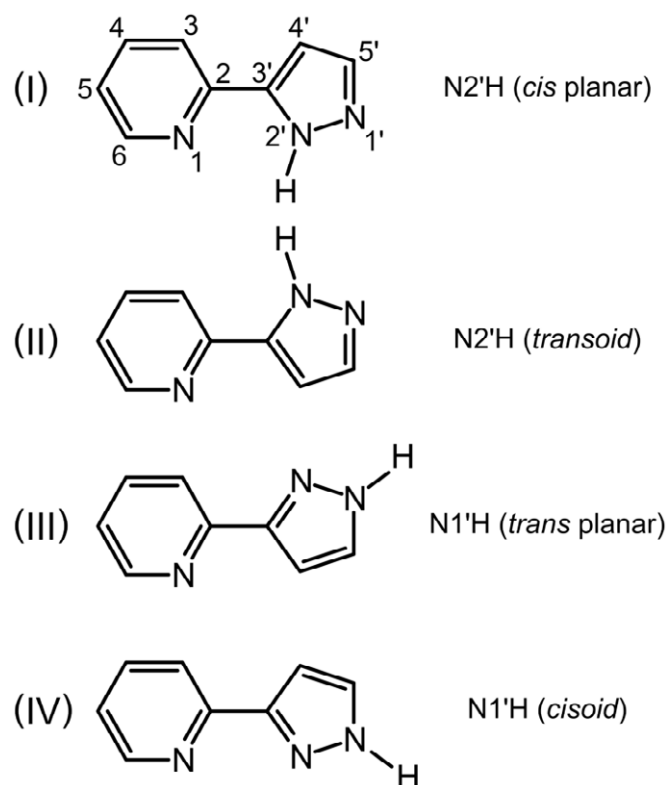
methods and refined by full-matrix least-squares methods based on F^2 using the SHELXL-97 and WinGX programs.^{3,4} All non-hydrogen atoms were refined anisotropically, and all the hydrogen atoms were included in geometrically calculated positions and refined with isotropic displacement parameters according to the riding model.

[1] Z. Otwinowsky, W. Minor, "DENZO and SCALEPACK". *Methods Enzymol.*1997, **276**, 307.

[2]CrysAlisPro, Oxford Diffraction (Poland), 2009.

[3]G. M. Sheldrick, SHELX97, "Programs for Crystal Structure Analysis; University of Göttingen". Göttingen, Germany, 1998.

[4]L. J. Farrugia, *J. Appl. Crystallogr.*1999, **32**, 837.



Scheme S-1 Hpyzpy in its four most stable tautomeric/rotameric forms: I > II ~ III > IV (according to ref. 2e). Two less stable forms (with proton at N1) are not shown.

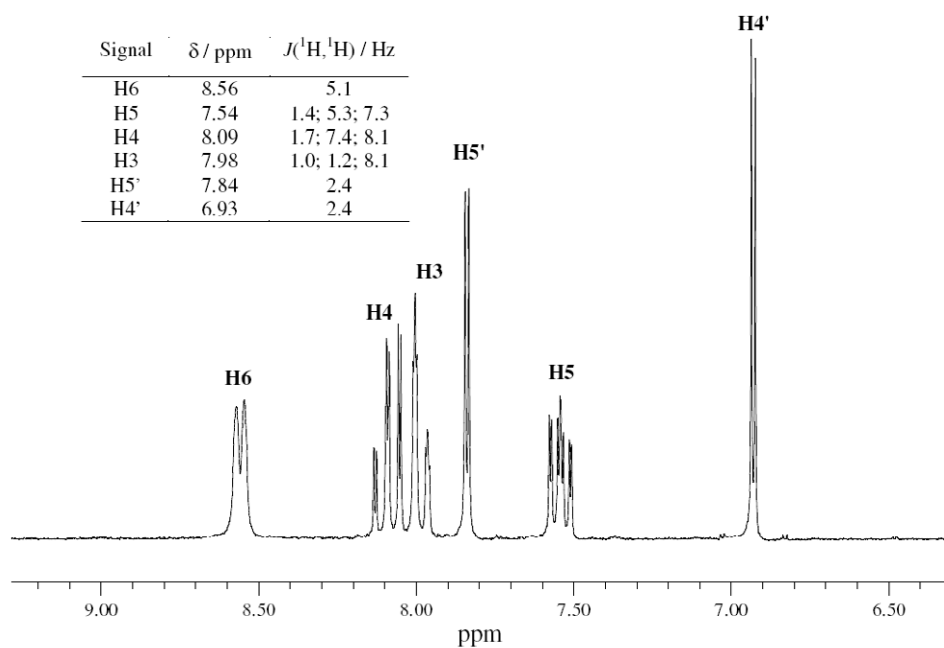


Figure S-1 ^1H NMR spectrum of Hpyzpy in D_2O , pH 6.1.

Table S-1 Selected distances (\AA) and angles ($^\circ$) for complex **1**.

Pt1–C11	2.284(4)	N2'-Pt1-N1	80.4(5)
Pt1–C12	2.287(4)	N2'-Pt1-C11	93.3(4)
Pt1–N1	2.017(13)	N1-Pt1-C12	94.6(4)
Pt1–N2'	1.989(12)	C11-Pt1-C12	91.68(16)
		N1-Pt1-C11	173.6(4)
		N2'-Pt1-C12	174.5(4)

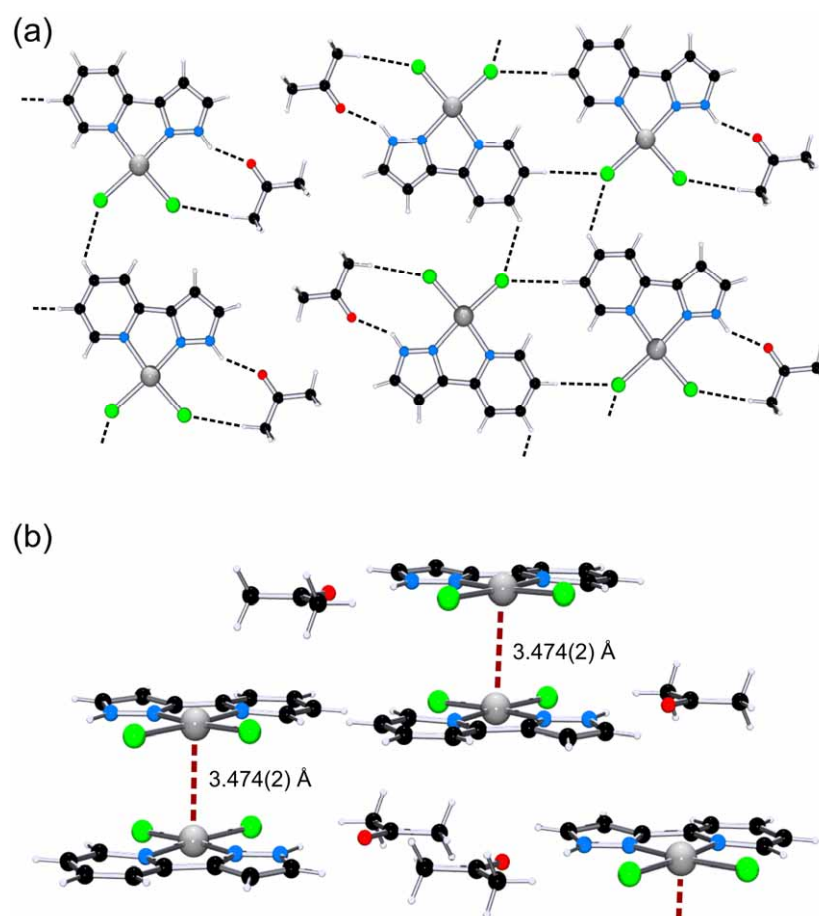


Figure S-2 (a) View of the hydrogen bonding pattern in **1**; (b) Stacking of the Pt(Hpyzpy-*N1,N2'*)Cl₂ units, with short Pt...Pt distances.

Table S-2 Selected distances (Å) and angles (°) for complex **2**.

Pt1–N1	2.033(4)	N2'–Pt1–N1	79.41(17)
Pt1–N2'	1.996(4)	N2'–Pt1–N1	100.59(17)
Pt2–N1a	2.033(4)	N1–Pt1–N1	180
Pt2–N1b	2.032(4)	N2'–Pt1–N2'	180
Pt2–N2a'	1.994(4)	N2a'–Pt2–N1a	79.91(16)
Pt2–N2b'	2.001(4)	N2a'–Pt2–N1b	100.07(16)
Pt1...Pt2	3.2985(2)	N2b'–Pt2–N1b	80.05(16)
		N1b–Pt2–N1a	178.60(16)
		N2a'–Pt2–N2b'	176.23(16)

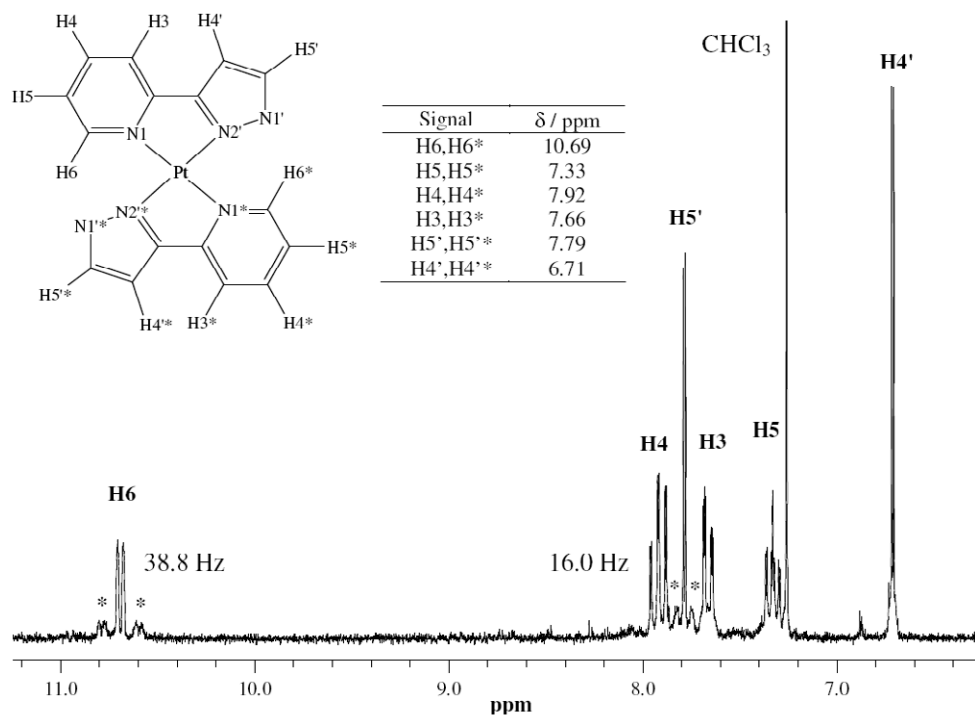


Figure S-3 ¹H NMR spectrum of **2** in CDCl₃.