# metal-organic papers

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#### Key indicators

Single-crystal X-ray study T = 296 KMean  $\sigma(C-C) = 0.020 \text{ Å}$  R factor = 0.058 wR factor = 0.174 Data-to-parameter ratio = 13.0

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# *µ*-Guanidinido-bis[(terpyridine)platinum(II)] tris(hexafluorophosphate) acetonitrile solvate

The title cationic complex  $\mu$ -guanidinido-bis[(2,2':6',2''-terpyridine)platinum(II)], [Pt<sub>2</sub>(C<sub>15</sub>H<sub>11</sub>N<sub>3</sub>)<sub>2</sub>(CH<sub>4</sub>N<sub>3</sub>)]<sup>3+</sup>, has a twofold axis and shows a stacked dinuclear structure with a short Pt···Pt distance of 3.0749 (6) Å. The complexes are further stacked to form a columnar structure in the crystal of the title compound. Received 8 April 2002 Accepted 23 April 2002 Online 11 May 2002

## Comment

The complex cation,  $[Pt_2(gua)(tpy)_2]^{3+}$  (tpy is 2,2':6',"-terpyridine and gua is guanidinide), has a stacked dinuclear structure of the Pt-tpy moiety bridged by a guanidine anion (Fig. 1), (I).



There exists a crystallographic twofold axis through the C16-N5 bond of the guanidinide ligand. The Pt-tpy moiety is almost planar and the deviations of the atoms are within 0.13 Å of the least-squares plane. The two Pt-tpy planes in the dinuclear complex are not parallel, but slightly tilted, with a dihedral angle of  $16.4 (1)^\circ$ , clearly because of the repulsion between the  $\pi$  systems of tpy. The Pt. Pt distance is 3.0749 (6) Å, which is short for divalent platinum, enabling the metal atoms to interact with each other electronically. In fact, the complex exhibits a very intense luminescence at around 630 nm, originating from the <sup>3</sup>MMLCT (metal-metal-to-ligand charge transfer) state, both in solution and in the solid state, even at room temperature. The structure is essentially the same as that of the corresponding perchlorate salt (Yip et al., 1992), although the latter has a lower symmetry than that of the  $PF_6^-$  salt. As shown in Fig. 2, the dinuclear complexes are stacked to form a columnar structure along the c axis. The adjacent dinuclear complexes in the column are related by an inversion center, and thus the adjacent Pt-tpy planes are completely parallel to each other, with an interplanar spacing of 3.48 (1) Å. The Pt atoms in adjacent complexes in the column are separated by a distance of 4.725 (1) Å. The acetonitrile solvate molecule is linked to the guanidinide ligand by a hydrogen bond.

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#### Figure 1

A perspective view of the title cation, showing the atom-numbering scheme, with displacement ellipsoids plotted at the 30% probability level.



# **Experimental**

[PtCl(tpy)]Cl was prepared according to the literature method (Howe-Grant & Lippard, 1980). The complex cation  $[Pt_2(gua)(tpy)_2]^{3+}$  was synthesized in a similar way to that reported by Yip et al. (1992). A mixture of [PtCl(tpy)]Cl·2H<sub>2</sub>O (54 mg, 0.1 mmol) and guanidine carbonate (9 mg, 0.05 mmol) in water was refluxed for 24 h. An excess of NH<sub>4</sub>PF<sub>6</sub> was added to the red solution. The resulting red precipitate was recrystallized from acetonitrile/ethanol to give red crystals of (I).

### Crystal data

[Pt <sub>2</sub> (C <sub>15</sub> H <sub>11</sub> N <sub>3</sub> ) <sub>2</sub> (CH <sub>4</sub> N <sub>3</sub> )](PF <sub>6</sub> ) <sub>3</sub>	$D_x = 2.087 \text{ Mg m}^{-3}$
$2C_2H_3N$	Mo K $\alpha$ radiation
$M_r = 1431.78$	Cell parameters fro
Monoclinic, $C2/c$	reflections
a = 17.000 (5)  Å	$\theta = 14.9 - 15.0^{\circ}$
b = 20.351 (4)  Å	$\mu = 6.33 \text{ mm}^{-1}$
c = 13.915 (4)  Å	T = 296.2  K
$\beta = 108.87 \ (2)^{\circ}$	Plate, red
$V = 4555 (1) \text{ Å}^3$	$0.60 \times 0.50 \times 0.08$
$\mathbf{Z} = \mathbf{A}$	

#### Data collection

Rigaku AFC-7R diffractometer  $\omega$ –2 $\theta$  scans Absorption correction: analytical (de Meulenaer & Tompa, 1965)  $T_{\min} = 0.418, T_{\max} = 0.999$ 6172 measured reflections 5243 independent reflections 3897 reflections with  $I > 2\sigma(I)$ 

#### Refinement

Refinement on  $F^2$  $R[F^2 > 2\sigma(F^2)] = 0.058$ wR(F<sup>2</sup>) = 0.174 S=1.023897 reflections 299 parameters

 $\alpha$  radiation parameters from 24 ections 4.9-15.0  $.33 \text{ mm}^{-1}$ 96.2 K red  $0.50 \times 0.08 \text{ mm}$ 

 $R_{\rm int} = 0.038$  $\theta_{\rm max} = 27.5^{\circ}$  $h = -2 \rightarrow 22$  $k=0\to 26$  $l = -18 \rightarrow 17$ 3 standard reflections every 150 reflections intensity decay: 5.6%

H-atom parameters constrained  $w = 1/[\sigma^{\hat{2}}({F_o}^2) + (0.136P)^2]$ where  $P = (F_o$  $(2^{2} + 2F_{c}^{2})/3$  $(\Delta/\sigma)_{\rm max} = 0.002$  $\Delta \rho_{\rm max} = 2.69 \text{ e} \text{ Å}^{-3}$  $\Delta \rho_{\rm min} = -3.42 \text{ e} \text{ Å}^{-3}$ 

## Table 1

Figure 2

Selected geometric parameters (Å, °).

Pt1-N1	2.024 (7)	Pt1-N4	1.987 (7)
Pt1-N2	1.922 (7)	N4-C16	1.34 (1)
Pt1-N3	2.018 (9)	N5-C16	1.35 (2)
N1-Pt1-N2	82.0 (3)	N2-Pt1-N4	177.1 (4)
N1-Pt1-N3	162.1 (3)	N3-Pt1-N4	98.4 (4)
N1 - Pt1 - N4	99.4 (3)	$N4 - C16 - N4^{i}$	119 (1)
N2-Pt1-N3	80.3 (3)	N4-C16-N5	120.5 (6)
$Pt1 \cdots Pt1^{i}$	3.0749 (6)		

The stacking arrangement of the  $[Pt_2(gua)(tpy)_2]^{3+}$  complex cations.

Symmetry code: (i)  $1 - x, y, \frac{1}{2} - z$ .

# Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
N4-H13···F1	0.88	2.41	3.14(1)	140
$N5-H14\cdots N6^{ii}$	0.87	3.00	3.16 (2)	93
$N5-H14\cdots F8^{iii}$	0.87	2.65	3.21 (1)	123

Symmetry codes: (i) 1 - x,  $y, \frac{1}{2} - z$ ; (ii)  $\frac{3}{2} - x, \frac{1}{2} - y, 1 - z$ ; (iii)  $x - \frac{1}{2}, \frac{1}{2} + y, z$ .

The data collection was successfully performed by using a crystal in a sealed capillary. Atoms for the acetonitrile solvent molecule were refined isotropically, because their large displacement parameters suggested disorder, but the anisotropic ellipsoids were featureless,

showing no particular characteristics that might suggest the direction of the disorder or the existence of disordered sites.

Data collection: *Rigaku/AFC Diffractometer Control Software* (Rigaku, 1995); cell refinement: *Rigaku/AFC Diffractometer Control Software*; data reduction: *TEXSAN* (Molecular Structure Corporation and Rigaku, 2000); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *TEXSAN*.

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