

Terpyridine-Platinum(II) Acetylides Complexes Bearing Pendent Coordination Units

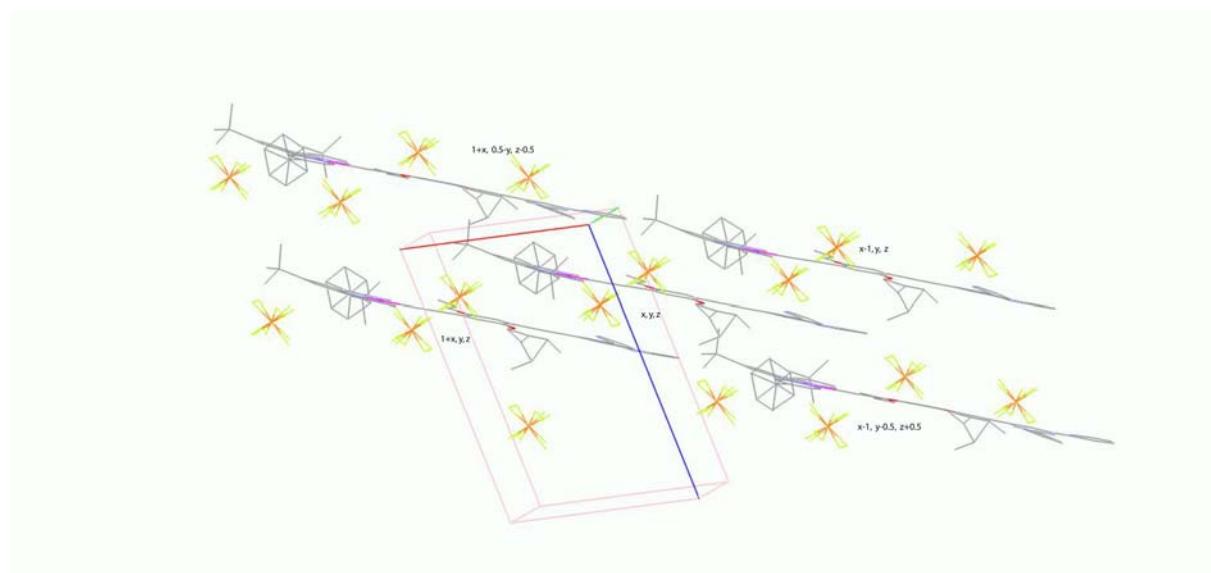
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Supporting Information (3 pages)

Table S1: π - π interactions for crystal compound 7

	Cg(I)	Cg(J)	[ARU(J)]	Cg-Cg (Å)	Dihedral angle (°)	Cg(I)_perp on ring J (Å)
Cycle Pt	Cg(1)	Cg(7)	1+X, 1/2-Y, -1/2+Z	3.889(3)	0.87	3.529
	Cg(2)	Cg(6)	1+X, 1/2-Y, -1/2+Z	3.614(4)	7.20	3.412
	Cg(3)	Cg(7)	1+X, 1/2-Y, -1/2+Z	3.962(4)	3.04	3.604
	Cg(4)	Cg(9)	1+X, Y, Z	3.872(4)	7.77	3.576
Cycle T	Cg(6)	Cg(2)	-1+X, 1/2-Y, 1/2+Z	3.614(4)	7.20	3.511
	Cg(7)	Cg(1)	-1+X, 1/2-Y, 1/2+Z	3.888(3)	0.87	3.524
	Cg(7)	Cg(3)	-1+X, 1/2-Y, 1/2+Z	3.962(4)	3.04	3.547
	Cg(8)	Cg(9)	-1+X, Y, Z	3.620(4)	3.33	3.457
Cycle B	Cg(9)	Cg(4)	-1+X, Y, Z	3.873(4)	7.77	3.647
	Cg(9)	Cg(8)	1+X, Y, Z	3.620(4)	3.33	3.505

Figure S1: packing of the crystal structure for compound 7



Captions for figure packing

in red, rings from the Pt moiety involved in π - π interactions

in green, central phenyl rings

in blue, rings from the t-pyridine moiety

For sake of clarity, main conformations in disorder were shown and PF6

in front of molecules in transparency.

(1) 1+X, 1/2-Y, -1/2+Z

(2) 1+X, Y, Z

(3) -1+X, 1/2-Y, 1/2+Z

(4) -1+X, Y, Z

Note: The elongation of the complex along its main dimension is slightly bent following the propeller geometry of the central phenyl ring (due to aliphatic chain di-substitution (C_n , $n=4$)) to let the concave space of the terpyridine moiety accommodate the (tert)butyl group of the neighbour molecule in the same plane (2+x, 1/2-y, z-1/2). (distance C53-C28 > distance C53-C38 (23.298 Å vs 22.124 Å and angle C53-Pt-N6 7.25° vs angle C53-Pt-N4 15.07°)