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Key indicators

Single-crystal X-ray study T = 298 KMean $\sigma(\text{C}-\text{C}) = 0.007 \text{ Å}$ Disorder in main residue R factor = 0.028 wR factor = 0.067 Data-to-parameter ratio = 16.3

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

cis-(2-Acetylcyclopentanonato- $\kappa^2 O, O'$)bis(2-pyridylphenyl- κN)iridium(III) dichloromethane solvate

The title compound, $[Ir(C_{11}H_8N)_2(C_7H_9O_2)]\cdot CH_2Cl_2$ or $[Ir(ppy)_2(2\text{-acetylcyclopentanone})]$ (ppy = 2-phenylpyridine), is a neutral mononuclear iridium(III) complex. The complex and solvent molecules both lie on twofold rotation axes. The iridium centre is coordinated by two N atoms and two C atoms from two ppy ligands [Ir-N = 2.044 (3) Å and Ir-C = 1.992 (4) Å], and two O atoms from the β -diketonate ligand [Ir-O= 2.11 (2) and 2.21 (2) Å], forming a distorted octahedral environment. The structure is stabilized by aromatic π - π stacking and C-H··· π interactions.

Comment

This work is related to a previously reported communication on the Ir^{III} complex with the bis(ppy) analogue, *cis*-bis-(phenylpyridine)(3-methylpentane-2,4-dionate)iridium(III) (Kim & Lee, 2006). A great number of cationic and neutral Ir^{III} complexes of the types [Ir(CN)₂ L_2]X and [Ir(CN)₂(LL)], where *CN* represents typically cyclometallating phenylpyridine derivatives, have been reported (Baldo *et al.*, 2000; Pomestchenko *et al.*, 2003; Duan *et al.*, 2003); *L* and *LL* are mono- and bidentate chelating anions, respectively, and X is any counter-anion. The title complex, (I), belongs to the latter type, where *CN* is ppy, and *LL* is 2-acetylcyclopentanone. The synthesis and electroluminescent properties of (I) have recently been reported by one of the authors and others (Kim, Paik *et al.*, 2004; Kim, Kim *et al.*, 2004).



Compound (I) (Fig. 1) is a solvated neutral mononuclear $[Ir(ppy)_2(2\text{-}acetylcyclopentanone)]$ (ppy is 2-phenylpyridine) complex. The asymmetric unit consists of one half-molecule of the complex and one half-molecule of solvent. Both molecules lie on twofold rotation axes. Bond distances for the coordination of Ir are summarized in Table 1. The Ir atom has a distorted octahedral geometry involving two ppy ligands and a β -diketonate ligand. The dihedral angle made by the N1/C1–C5 ring and the C6–C11 ring of ppy is 2.6 (3)°.

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The crystal structure is stabilized by aromatic π - π stacking and Cl₂HC-H··· π interactions (Fig. 2). The shortest parallel ring centroid separation is $Cg1 \cdots Cg2^{ii}$ of 3.748 (7) Å and the shortest distance is $Cg2\cdots$ H19(A,B)-C19^v of 2.60 Å [Cg1 and Cg2 are the centroids of the C1-C5/N and C6-C11 rings; symmetry codes as in Fig. 2].

Experimental

A mixture of [IrCl(ppy)₂IrCl]₂(83.6 mg, 0.078 mmol), 2-acetylcyclopentanone (25.2 mg, 0.2 mmol) and potassium carbonate (82.9 mg, 0.6 mmol) in 2-ethoxyethanol (5 ml) was stirred under reflux for 1 h. After cooling the reaction mixture to room temperature, the yellow precipitate was filtered off, and washed with water, followed by two portions of methanol (5 ml). The crude product was separated by chromatography on silica gel with dichloromethane as eluent to give a yellow solid (71.3 mg, 73%). Single crystals suitable for X-ray diffraction were grown by slow diffusion of hexane into a dichloromethane solution.

Crystal data

$[Ir(C_{11}H_8N)_2(C_7H_9O_2)] \cdot CH_2Cl_2$
$M_r = 710.64$
Monoclinic, $C2/c$
a = 12.094 (1) Å
b = 14.889 (2) Å
c = 15.287 (2) Å
$\beta = 96.743 \ (9)^{\circ}$
V = 2733.7 (6) Å ³

Data collection

Stoe STADI-4 diffractometer $\omega - 2\theta$ scans Absorption correction: numerical (X-SHAPE; Stoe & Cie, 1996) $T_{\min} = 0.431, \ T_{\max} = 0.533$ 3137 measured reflections

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0313H)]$
$R[F^2 > 2\sigma(F^2)] = 0.028$	+ 8.9226P]
$wR(F^2) = 0.067$	where $P = (F_0^2 + 2F_c^2)$
S = 1.07	$(\Delta/\sigma)_{\rm max} < 0.001$
3137 reflections	$\Delta \rho_{\rm max} = 0.70 \ {\rm e} \ {\rm \AA}^{-3}$
192 parameters	$\Delta \rho_{\rm min} = -0.53 \ {\rm e} \ {\rm \AA}^{-3}$
H-atom parameters constrained	

Table 1

Selected bond lengths (Å).

Ir1-C11	1.992 (4)	C12-C16	1.468 (8)
Ir1-N1	2.044 (3)	C12-C13	1.500 (9)
Ir1-O1	2.11 (2)	C13-C14	1.478 (8)
Ir1-O2	2.21 (2)	C14-C15	1.479 (9)
O1-C12	1.258 (7)	C15-C16	1.507 (7)
O2-C17	1.269 (8)	C16-C17	1.358 (8)

The 2-acetylcyclopentanone ligand is disordered over a twofold rotation axis and all non-H atoms were restrained to lie in a plane with half-occupancy. Bond distances were restrained to acceptable values; the displacement ellipsoids of the pairs O1/O2, C12/C17 and C13/C18 were set equal to each other. The vibration of C16 was restrained to be nearly isotropic. All H atoms were positioned geometrically and refined using a riding model, with C-H = 0.93 Å for aromatic, 0.97 Å for methylene, and 0.96 Å for methyl H atoms,

Z = 4 $D_r = 1.727 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation $\mu = 5.11 \text{ mm}^{-1}$ T = 298 (2) K Block, yellow $0.50 \times 0.25 \times 0.20$ mm

3137 independent reflections 2873 reflections with $I > 2\sigma(I)$ $\theta_{\rm max} = 27.5^{\circ}$ 3 standard reflections frequency: 60 min intensity decay: 3.2%

P)² /3



Figure 1

The molecular structure of (I), shown with 30% probability displacement ellipsoids. One disordered component of 2-acetylcyclopentanone and the solvent (CH₂Cl₂) have been omitted. [Symmetry code: (i) 1 - x, $y, \frac{3}{2} - z$.]



Figure 2

The π - π and Cl₂HC-H··· π interactions in (I), shown as red and blue dotted lines, respectively. Cg1 and Cg2 are the centroids of the C1-C4/N and C6-C11 rings, respectively. H atoms of ppy have been omitted for clarity. [Symmetry codes: (i) $1 - x, y, \frac{3}{2} - z$; (ii) 1 - x, -y, 1 - z; (iii) x, -y, -y, 1 - z; (iii) x, -y, -y, -y, -z; (iii) x, -y, -y, -z; (iii) x, -y, -y, -z; (iii) x, $\frac{1}{2} + z$; (iv) 1 - x, -y, 2 - z; (v) $\frac{1}{2} + x, y - \frac{1}{2}, z$.]

and with $U_{iso}(H) = 1.2U_{eq}(C)$ for aromatic and methylene H atoms, and $U_{iso}(H) = 1.5U_{eq}(C)$ for methyl H atoms.

Data collection: STADI4 (Stoe & Cie, 1996); cell refinement: STADI4; data reduction: X-RED32 (Stoe & Cie, 1996); program(s) used to solve structure: SHELXS97-2 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97-2 (Sheldrick, 1997); molecular graphics: ORTEP-3 (Farrugia, 1997) and DIAMOND (Brandenburg, 1998); software used to prepare material for publication: SHELXL97-2.

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