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Key indicators

Single-crystal X-ray study
 $T = 298\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.008\text{ \AA}$
 R factor = 0.030
 wR factor = 0.076
Data-to-parameter ratio = 19.0For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.An unexpected product: di- μ -iodo-bis-
{[2-(*N*-diphenylphosphinomethyl-*N*-cyclo-
hexylamino)pyridine]iodomercury(II)}

The title compound, $[\text{Hg}_2\text{I}_4(\text{C}_{24}\text{H}_{27}\text{N}_2\text{P})_2]$, is found to be a symmetric doubly iodide-bridged dimer, with the phosphine ligands attached *trans* to the two Hg atoms [$\text{Hg}-\text{P} = 2.4726(12)\text{ \AA}$]. The molecule has a centre of symmetry at the centre of the four-membered ring formed by the two Hg and two bridging I atoms.

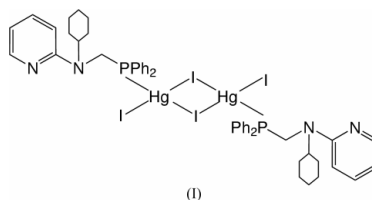
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Comment

Pyridylphosphines continue to generate much interest as excellent ligands for stabilizing many transition-metal coordination and organometallic complexes (Espinete & Soulantica, 1999). One important property of these ligands is that they can stabilize metal ions in a variety of valence states and geometries. Hence, a metal-metal bond between an electron-rich metal (soft base) and a high-oxidation-state metal (Lewis acid) is easily formed (Zhang & Cheng, 1996). Recently, we designed and synthesized a new pyridylphosphine ligand, namely 2-(*N*-diphenylphosphinomethyl-*N*-cyclohexylamino)-pyridine, *L*, and obtained a hetero-binuclear complex containing an Fe-Cu bond (Cui *et al.*, 2001). However, when *trans*- $\text{Fe}(\text{CO})_3\text{L}_2$ was reacted with HgI_2 , instead of the desired compound containing an Fe-Hg bond, the title compound, (I), was unexpectedly formed.



In (I), each Hg atom is four-coordinated by two bridging I atoms (Fig. 1), [$\text{Hg1}-\text{I1} 2.8036(4)\text{ \AA}$ and $\text{Hg1}-\text{I1}^i 2.9667(4)\text{ \AA}$], one terminal I atom [$\text{Hg1}-\text{I2} 2.7699(4)\text{ \AA}$] and one P atom from the ligand *L* [$\text{Hg1}-\text{P4} 2.4723(11)\text{ \AA}$]. There is significant distortion from tetrahedral geometry, the angles about the metal ranging from $94.047(10)$ to $127.87(3)^\circ$ (Table 1).

The title compound may be compared with the series $(\text{PPh}_3)_2\text{Hg}_2\text{X}_4$ which have the same molecular structure as (I) ($X = \text{Cl}, \text{Br}, \text{I}$), but with replacement of ligand *L* by triphenylphosphine (Bell *et al.*, 1980, 1989*b*; Bowmaker *et al.*, 1993). As in (I), the complex with $X = \text{Cl}$ is centrosymmetric, whereas the bromo and iodo complexes are non-centrosymmetric. While the larger $X_{\text{br}}-\text{Hg}-X_{\text{br}}$ angles (X_{br} is bridging halogen) and $\text{Hg}-X$ distances in the iodide complex may be readily attributed to the increasing size of the halogen, no

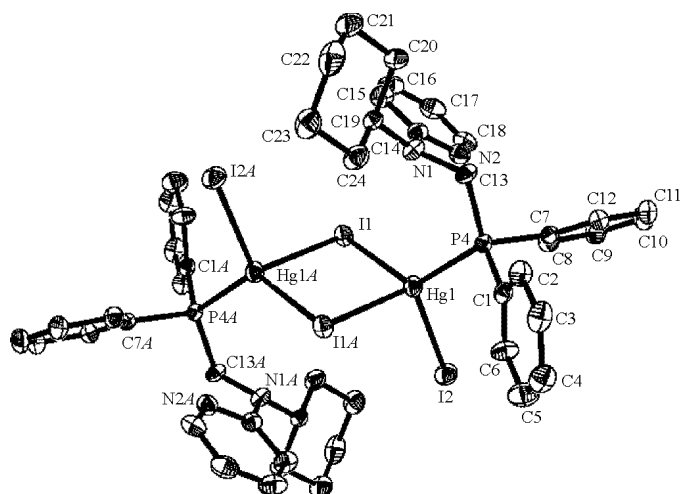


Figure 1
A view of (I), showing the atomic numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. Suffix *A* corresponds to symmetry code (i) in Table 1

ready explanation is available for the substantial asymmetry of the Hg— X_{br} distances found in the iodide complex. In contrast to $(R_3P)_2HgX_2$ complexes, in which the P—Hg—P angle is an effective indicator of the σ -donating ability of both the phosphine and the halogen (Allen *et al.*, 1985; Bell *et al.*, 1989a), the P—Hg— X_t angles (X_t is terminal halogen) in the present complex, which might, on the basis of the stronger σ -donor properties of iodide compared to chloride and bromide, have been anticipated to be larger than in the chloride and bromide, are, in fact, very similar. The relative σ -donating ability of the halides is, however, reflected in the Hg—P distance. Thus the significantly larger distance found in the iodide complex may be attributed to the greater σ -donor ability of iodide compared to chloride and bromide.

Experimental

The synthesis was carried out under an Ar atmosphere. To a solution of *trans*-Fe(CO) $_3$ L $_2$ (0.23 g, 0.25 mmol) was added solid HgI $_2$ (0.11 g, 0.25 mmol). After stirring for 5 min, the solid disappeared. The mixture was filtered and the solvent was removed. The residue was recrystallized from CH $_2$ Cl $_2$ /CH $_3$ OH and gave 0.17 g (85%) of a dark yellow solid.

Crystal data

[Hg $_2$ I $_4$ (C $_{24}$ H $_{27}$ N $_2$ P) $_2$]	$D_x = 2.170 \text{ Mg m}^{-3}$
$M_r = 1657.67$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 634 reflections
$a = 12.5386 (7) \text{ \AA}$	$\theta = 2.8\text{--}25.3^\circ$
$b = 16.1113 (9) \text{ \AA}$	$\mu = 8.58 \text{ mm}^{-1}$
$c = 12.6488 (7) \text{ \AA}$	$T = 298 (2) \text{ K}$
$\beta = 96.773 (1)^\circ$	Plate, dark yellow
$V = 2537.4 (2) \text{ \AA}^3$	$0.40 \times 0.35 \times 0.05 \text{ mm}$
$Z = 2$	

Data collection

Bruker SMART CCD area-detector diffractometer	5142 independent reflections
φ and ω scans	4365 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	$R_{\text{int}} = 0.041$
11685 measured reflections	$\theta_{\text{max}} = 26.4^\circ$
	$h = -15 \rightarrow 15$
	$k = -18 \rightarrow 20$
	$l = -11 \rightarrow 15$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0458P)^2 + 0.249P]$
$R[F^2 > 2\sigma(F^2)] = 0.030$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.076$	$(\Delta/\sigma)_{\text{max}} = 0.002$
$S = 0.98$	$\Delta\rho_{\text{max}} = 1.49 \text{ e \AA}^{-3}$
5142 reflections	$\Delta\rho_{\text{min}} = -1.15 \text{ e \AA}^{-3}$
271 parameters	
H-atom parameters constrained	

Table 1

Selected geometric parameters (\AA , $^\circ$).

Hg1—P4	2.4723 (11)	Hg1—I1	2.8036 (4)
Hg1—I2	2.7699 (4)	Hg1—I1 ⁱ	2.9667 (4)
P4—Hg1—I2	115.61 (3)	I2—Hg1—I1 ⁱ	105.587 (13)
P4—Hg1—I1	127.87 (3)	I1—Hg1—I1 ⁱ	94.047 (10)
I2—Hg1—I1	105.332 (13)	Hg1—I1—Hg1 ⁱ	85.953 (10)
P4—Hg1—I1 ⁱ	104.03 (3)		

Symmetry code: (i) $2 - x, 2 - y, 1 - z$.

All H atoms were positioned geometrically, with C—H = 0.96 \AA . They were treated as riding atoms, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. The maximum and minimum electron-density peaks are located 0.92 and 0.88 \AA from Hg1, respectively.

Data collection: SMART (Bruker, 1998); cell refinement: SMART; data reduction: SAINT (Bruker, 1998) and SHELXTL (Bruker, 1997); program(s) used to solve structure: SHELXTL; program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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