metal-organic compounds

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3,5-Bis{4-[(benzimidazol-1-yl)methyl]phenyl}-4H-1,2,4-triazol-4-amine and its one-dimensional polymeric complex with HgCl₂

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The molecule of 3,5-bis{4-[(benzimidazol-1-yl)methyl]phenyl}-4*H*-1,2,4-triazol-4-amine (*L*), $C_{30}H_{24}N_8$, has an antiperiplanar conformation of the two terminal benzimidazole groups and forms two-dimensional networks along the crystallographic *b* axis *via* two types of intermolecular hydrogen bonds. However, in *catena*-poly[[[dichloridomercury(II)]- μ -3,5-bis{4-[(benzimidazol-1-yl)methyl]phenyl}-4*H*-1,2,4-triazol-4-amine] dichloromethane hemisolvate], {[HgCl₂(C₃₀H₂₄N₈)]·0.5CH₂-Cl₂)_n, synthesized by the combination of *L* with HgCl₂, the *L* ligand adopts a synperiplanar conformation. The Hg^{II} cation lies in a distorted tetrahedral environment, which is defined by two N atoms and two Cl atoms to form a one-dimensional zigzag chain. These zigzag chains stack *via* hydrogen bonds which expand the dimensionality of the structure from one to two.

Comment

Organometallic complexes have attracted considerable attention because of their fascinating topological structures (Eddaoudi *et al.*, 2002) and potential applications as functional materials in gas storage (Ma & Zhou, 2010), host–guest chemistry (Yoshizawa *et al.*, 2006), catalysis (Kim *et al.*, 2010) and luminescence (Cui *et al.*, 2012). Over the past decade, the design and construction of rigid organic ligands bridged by 1,2,4-triazol-4-amine has been pursued, due to their diversity in coordination chemistry and model applications in functional materials (Wang *et al.*, 2007, 2009; Liu *et al.*, 2009, 2010).

To date, various organic ligands have been used as molecular building blocks, but the use of 1,2,4-triazol-4-aminebased bent organic ligands as semi-rigid components has remained rare until recently. In order to investigate how the semi-rigid organic ligands bridged by 1,2,4-triazol-4-amine affect the arrangement of molecular complexes in selfassembled aggregates, we synthesized a new 1,2,4-triazol-4amine bridging ligand, namely 3,5-bis{4-[(benzimidazol-1-yl)- methyl]phenyl}-4*H*-1,2,4-triazol-4-amine, denoted *L* or (I). This compound introduces two large aromatic benzimidazole groups to the ligand. The combination of (I) with HgCl₂ afforded { $[HgCl_2L]\cdot CH_2Cl_2$ _n, (II), which features hydrogenbonded stacking-driven two-dimensional networks.



Within the free ligand, (I) (Fig. 1), the terminal benzimidazole groups adopt an antiperiplanar conformation across the central bridging 1,2,4-triazol-4-amine group. The dihedral angles between the planes of the benzimidazole and triazole rings are 58.1 (2) and 88.06 (18)° for the benzimidazole rings containing atoms N5 and N7, respectively, and the dihedral angles between the benzimidazole and adjacent benzene rings are 78.64 (18) and 80.37 (18)°, respectively. The two benzene rings of (I) are almost coplanar, with a dihedral angle of 7.3 (2)°.

In the crystal structure, molecules of (I) are arranged in chains *via* N4–H4A···N8ⁱ hydrogen bonds along the [101] direction (details and symmetry codes are given in Table 1). The chains stack *via* a second hydrogen-bond interaction (N4–H4B···N1ⁱⁱ) to form a two-dimensional network which





The molecular structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.



Figure 2

The two-dimensional structure of (I), constructed by hydrogen bonds (dashed lines). Uninvolved H atoms have been omitted for clarity. [Symmetry codes: (i) $x - \frac{1}{2}$, $-y + \frac{1}{2}$, $z - \frac{1}{2}$; (ii) x - 1, y, z.]

lies parallel to the (010) plane (Fig. 2). This is in contrast with 3,5-bis(2-chlorophenyl)-1H-1,2,4-triazol-4-amine (Zachara *et al.*, 2004), the molecules of which are linked by hydrogen bonds to form a one-dimensional chain.

Compound (II) crystallizes with one unique four-coordinated Hg^{II} centre in a distorted tetrahedral {HgCl₂N₂} environment involving two Cl atoms (Cl1 and Cl2) and two N atoms [N2 and N6ⁱ; symmetry code: (i) $x, -y + \frac{3}{2}, z - \frac{1}{2}$] from two different ligands (Fig. 3 and Table 2). Neighbouring Hg^{II} cations are bound together by triazole atom N2 and the terminal benzimidazole N6 atom of the *L* ligand to form an {HgCl₂*L*}_{*n*} one-dimensional zigzag chain. The coordination behaviour of the Hg^{II} cation is similar to that observed in Hg[1-(pyridin-2-ylmethyl)-1*H*-benzotriazole]Cl₂ (Liu *et al.*, 2008), where the 1-(pyridin-2-ylmethyl)-1*H*-benzotriazole ligand (L2) is also coordinated to two Hg^{II} centres to form a single chain. Similarly, in [Hg(L2)Cl₂], there are no bonding interactions observed between adjacent chains.

The ligand *L* adopts different conformations under different conditions. In the solid state, the benzimidazole moieties adopt a synperiplanar orientation about the triazole core in the free ligand, whereas after coordination to Hg^{II} the benzimidazole groups adopt a *cis* conformation. Compared with those given above for (I), the dihedral angles in (II) between the planes of the benzimidazole groups and the adjacent benzene rings change to 87.9 (3) and 75.8 (3)° for the benzimidazole rings containing atoms N5 and N7, respectively. These are nearly perpendicular, clearly as a result of coordination to Hg^{II}. Additionally, the dihedral angle formed by the two benzene rings changes from 7.3 (2)° in the free ligand to 18.9 (3)° in (II).

In the solid state, the zigzag chains in (II) are arranged along the *c* axis, where they interact *via* N4-H4A···Cl2ⁱⁱⁱ hydrogen-bond interactions, as shown in Fig. 4 (details in Table 3; symmetry code given in Table 3). The result is that a two-dimensional sheet is generated in the *bc* plane. The dichloromethane solvent molecules are located within the cavities formed by the layered stacking of (II) (Fig. 5), although it appears that only about half of the solvent molecule sites are occupied.

In summary, a new compound with a common zigzag chain motif has been successfully obtained based on a new 1,2,4triazol-4-amine bridging bent organic ligand, (I), and HgCl₂. The chains assemble through hydrogen bonds to form a twodimensional network. The hydrogen-bond interactions play an important role in constructing high-dimensional supramolecular compounds.



Figure 3

The molecular structure of (II), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. [Symmetry code: (i) x, $-y + \frac{3}{2}$, $z - \frac{1}{2}$.]



Figure 4

The two-dimensional structure of (II), constructed by hydrogen bonds (dashed lines). Uninvolved H atoms have been omitted for clarity. [Symmetry codes: (ii) x, $-y + \frac{3}{2}$, $z + \frac{1}{2}$; (iii) x, y - 1, z.]

Experimental

For the preparation of (I), 80% hydrazine hydrate (2.8 g, 45 mmol) was added with stirring to a solution of 3,5-bis{4-[(benzimidazol-1yl)methyl]phenyl]-1,3,4-oxadiazole (1.49 g, 3.0 mmol) in N,N-dimethylformamide (DMF; 20 ml). The mixture was stirred for 4 h at 423 K, then cooled to room temperature and poured into water (100 ml). The product was obtained as a white solid and purified on a silica-gel column using tetrahydrofuran as the eluent to afford (I) as a white crystalline solid (yield 1.00 g, 67%). A solution of (I) (5.00 mg, 0.010 mmol) in CH₂Cl₂ (10 ml) was left for about 2 d at room temperature, after which time colourless crystals were obtained (yield 3.24 mg, 65%). IR (KBr pellet, ν , cm⁻¹): 3442 (s), 1633 (s), 1496 (s), 1385 (s), 1295 (m), 1200 (w), 1123 (w), 974 (w), 765 (s), 619 (m); ¹H NMR (300 MHz, DMSO, 298 K, TMS): δ 8.44 (s, 2H, -C₃H₂N₂-), 7.96-7.94, 7.47-7.44 (aabb, 8H, -C₆H₄-), 7.67-7.64 (d, 2H, -C₃H₂N₂-, 7.56-7.53 (d, 2H, -C₆H₃-), 7.23-7.18 (m, 4H, -C₆H₃-), 6.17 (s, 2H, -NH₂), 5.57 (s, 4H, -CH₂-). Elemental analysis calculated for C₃₀H₂₄N₈: C 72.32, H 4.91, N 22.77%; found: C 72.56, H 4.87, N 22.57%.

For the synthesis of (II), a solution of $HgCl_2$ (5.42 mg, 0.020 mmol) in CH₃OH (5 ml) was layered onto a solution of (I) (9.93 mg,

0.020 mmol) in CH₂Cl₂ (8 ml). The mixture was left for about a week at room temperature and colourless crystals of (II) were obtained (yield 8.64 mg, 62%). IR (KBr pellet, v, cm⁻¹): 3447 (w), 1636 (s), 1509 (w), 1459 (m), 1384 (w), 1266 (m), 1193 (m), 740 (s). Elemental analysis calculated for C₆₁H₅₀Cl₆Hg₂N₁₆: C 45.34, H 3.23, N 13.66%; found: C 45.20, H 3.11, N 13.82%.

Compound (I)

Crystal data

 $C_{30}H_{24}N_8$ $M_r = 496.57$ Monoclinic, *Cc* a = 6.141 (2) Å b = 19.914 (6) Å c = 19.910 (6) Å $\beta = 96.537$ (6)°

Data collection

Bruker SMART CCD area-detector diffractometer Absorption correction: multi-scan (*SADABS*; Bruker, 2003) $T_{\rm min} = 0.986, T_{\rm max} = 0.992$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.053$ $wR(F^2) = 0.116$ S = 1.002748 reflections 343 parameters

Compound (II)

Crystal data

[HgCl₂($C_{30}H_{24}N_8$)]-0.5CH₂Cl₂ $M_r = 810.53$ Monoclinic, $P2_1/c$ a = 16.627 (3) Å b = 10.6199 (18) Å c = 20.082 (3) Å $\beta = 113.586$ (2)°

Data collection

- Bruker SMART CCD area-detector diffractometer
- Absorption correction: multi-scan (SADABS; Bruker, 2003) $T_{min} = 0.239, T_{max} = 0.280$

 $V = 2418.8 (13) \text{ Å}^{3}$ Z = 4Mo K\alpha radiation $\mu = 0.09 \text{ mm}^{-1}$ T = 298 K $0.17 \times 0.11 \times 0.09 \text{ mm}$

5790 measured reflections 2748 independent reflections 2008 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.063$

2 restraints H-atom parameters constrained $\Delta \rho_{max} = 0.14 \text{ e } \text{ Å}^{-3}$ $\Delta \rho_{min} = -0.15 \text{ e } \text{ Å}^{-3}$

 $V = 3249.9 (10) Å^{3}$ Z = 4 Mo K\alpha radiation \mu = 5.02 mm^{-1} T = 298 K 0.40 \times 0.38 \times 0.34 mm

16724 measured reflections 6097 independent reflections 4599 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.040$



Figure 5

The crystal packing of (II), viewed along the b axis. The dichloromethane solvent molecules are located in channels.

Table 1	
Hydrogen-bond geometry (Å, °) for (I).	

$D-\mathrm{H}\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
N4–H4A····N8 ⁱ	0.89	2.53	3.354 (5)	155
$N4-H4B\cdots N1^{ii}$	0.89	2.50	3.222 (6)	138

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

Table 2

Selected geometric parameters (Å, °) for (II).

Cl1-Hg1	2.4331 (17)	$Hg1-N6^{i}$	2.265 (4)
Cl2-Hg1	2.3793 (17)	Hg1-N2	2.353 (4)
N6'-Hg1-N2	95.32 (15)	N6 ⁴ -Hg1-Cl1	109.17 (12)
N6 ⁱ -Hg1-Cl2	112.97 (12)	N2-Hg1-Cl1	103.99 (11)
N2-Hg1-Cl2	111.62 (13)	Cl2-Hg1-Cl1	120.51 (7)

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

Table 3

Hydrogen-bond geometry (Å, °) for (II).

$D-\mathrm{H}\cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
N4-H4 B ···N8 ⁱⁱ	0.89	2.09	2.940 (8)	160
N4-H4 A ···Cl2 ⁱⁱⁱ	0.89	2.98	3.473 (5)	117

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

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.038$	6 restraints
$wR(F^2) = 0.102$	H-atom parameters constrained
S = 1.04	$\Delta \rho_{\rm max} = 1.22 \text{ e} \text{ Å}^{-3}$
6097 reflections	$\Delta \rho_{\rm min} = -0.64 \text{ e } \text{\AA}^{-3}$
397 parameters	

The H atoms on N4 were located in a difference map and their positions were initially refined subject to an N—H distance restraint of 0.89 (2) Å. Subsequently, these H atoms were treated as riding atoms, with $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm N})$. The remaining H atoms were placed in geometrically idealized positions and included as riding atoms, with C—H = 0.93 Å and $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C})$ for aromatic H atoms or C—H = 0.97 Å and $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C})$ for methylene H atoms.

In noncentrosymmetric (I), Friedel pairs were merged in the final refinement and the absolute structure was chosen arbitrarily. In (II), the dichloromethane solvent molecule was refined with the C–Cl and Cl···Cl distances restrained to 1.78 (1) and 2.80 (2) Å, respectively, in order to retain reasonable geometry. The solvent sites also appeared to be only partially occupied and a fixed site-occupation factor of 0.5 was employed for chemical and crystallographic rationality. The main directions of movement of covalently bonded atoms C29, C30 and C31 were likewise restrained to be similar, with an s.u. value of 0.005 Å² (Müller *et al.*, 2006).

For both compounds, data collection: *SMART* (Bruker, 2003); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 2003); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: OV3013). Services for accessing these data are described at the back of the journal.

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