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$(\mu_2$ -Adipato- κ^4 O,O':O",O"')bis[aqua(benzene-1,2diamine- $\kappa^2 N$, N')chloridocadmium]: crystal structure and Hirshfeld surface analysis

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The full molecule of the binuclear title compound, $[\text{Cd}_2\text{Cl}_2(\text{C}_6\text{H}_8\text{O}_4)(\text{C}_6\text{H}_8\text{N}_2)_2$ - $(H₂O)₂$, is generated by the application of a centre of inversion located at the middle of the central CH_2-CH_2 bond of the adipate dianion; the latter chelates a Cd^H atom at each end. Along with two carboxylate-O atoms, the Cd^H ion is coordinated by the two N atoms of the chelating benzene-1,2-diamine ligand, a Cl^- anion and an aqua ligand to define a distorted octahedral $CdCN_2O_3$ coordination geometry with the monodentate ligands being mutually cis. The disparity in the Cd—N bond lengths is related to the relative trans effect exerted by the Cd—O bonds formed by the carboxylate-O and aqua-O atoms. The packing features water-O $-H \cdot \cdot \cdot O$ (carboxylate) and benzene-1,2-diamine-N $H \cdots$ Cl hydrogen bonds, leading to layers that stack along the *a*-axis direction. The lack of directional interactions between the layers is confirmed by a Hirshfeld surface analysis.

1. Chemical context

In the +II oxidation state, the $4d^{10}$ cadmium(II) cation is a favourite of researchers studying coordination polymers/ metal–organic frameworks. With the ability to readily coordinate a variety of different donor atoms, i.e. both hard and soft donors, and to adopt a range of coordination geometries, a diverse array of structures can be generated. The motivation for studying cadmium(II) compounds in this context, over and above intellectual curiosity, rests primarily with evaluating their photoluminescence properties (Lestari et al., 2014; Xue et al., 2015; Seco et al., 2017).

Our interest in cadmium(II) structural chemistry is in the controlled formation (dimensionality and topology) of coordination polymers of dithiophosphates $(\overline{S_2P(OR)_2};$ Lai & Tiekink, 2004, 2006), xanthates $($ ⁻S₂COR; Tan, Azizuddin et al., 2016) and dithiocarbamates $($ ⁻S₂CNR₂; Chai et al., 2003), in particular those substituted with hydroxyethyl groups, capable of forming hydrogen-bonding interactions (Tan et al.,

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2013; Tan, Halim & Tiekink, 2016). In this connection, we now describe the crystal structure determination and Hirshfeld surface analysis of a cadmium(II) species, (I), with a potentially bridging adipato dianion and an ancillary ligand, benzene-1,2-diamine, capable of forming hydrogen-bonding interactions.

2. Structural commentary

The asymmetric unit of (I) comprises half a molecule of (I), Fig. 1, with the full molecule generated about a centre of inversion. The key feature of the structure is the tetra-coordinate mode of coordination of the adipato dianion, linking the two Cd^H cations. Each carboxylate group forms equivalent Cd—O bonds, the difference in the two bonds being only 0.01 Å, Table 1. More asymmetry is found in the coordination of the benzene-1,2-diamine ligand with the Cd—N1 bond length being 0.05 Å longer than Cd—N2. This may be traced to the different trans effects exerted by the oxygen atoms in that the N1 atom is *trans* to the carboxylate-O1 atom $[N1]$ $Cd - O1 = 166.89 (6)°$] whereas N2 is opposite to the coordinating water molecule $[N2-Cd-O1W = 149.12 (7)°]$. The coordination geometry is completed by the chloride anion which, owing to the presence of two chelating ligands, occupies a position *cis* to the aqua group. The donor set is CIN_2O_3 and defines a distorted octahedral geometry.

As might be expected, the four-membered chelate ring formed by the carboxylate group is strictly planar (r.m.s. deviation = 0.0009 Å). There is a twist in the chain of the dicarboxylate ligand with the bond linking the quaternary atom to the aliphatic group being $+$ *anti-clinal, i.e.* the $O2-$ C1—C2—C3 torsion angle is 145.7 (3) $^{\circ}$ but, - anti-periplanar about the central bond, *i.e.* $C1 - C2 - C3 - C3^i$ is -177.6 (3)[°]; symmetry code: (i) $-x$, $2 - y$, $-z$. There is a distinct kink in the five-membered ring formed by the benzene-1,2-diamine ligand. This is readily seen in the dihedral angle of 58.57 $(7)^\circ$ formed between the plane through the $CdN₂$ atoms and the benzene ring.

Figure 1

The molecular structure of (I), showing the atom-labelling scheme and displacement ellipsoids at the 70% probability level. The molecule is disposed about a centre of inversion and unlabelled atoms are related by the symmetry operation $(-x, 2 - y, -z)$.

Table 2 Hydrogen-bond geometry (\AA, \degree) .

 $2.398(2)$
 $2.5283(6)$

 $2.2265(18)$

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

3. Supramolecular features

As summarized in Table 2, all acidic hydrogen atoms in the molecule of (I) are involved in conventional hydrogenbonding interactions. The water-H atoms each form an hydrogen bond with a carboxylate-O atom to form strands

Molecular packing in (I) : (a) a view of the supramolecular layer parallel to (100) sustained by water-O $-H \cdot \cdot \cdot O(carboxulate)$ and benzene-1,2diamine-N-H \cdots Cl hydrogen bonds and (b) a view of the unit-cell contents in projection down the b axis. The O $-H$ \cdots O and N $-H$ \cdots Cl hydrogen bonds are shown as orange and blue dashed lines, respectively.

Figure 3

A view of the Hirshfeld surface for (I) mapped over d_{norm} in the range -0.597 to +1. 425 au.

propagating along the b-axis direction, involving the carboxylate-O1 atoms, and along the c-axis direction, involving the carboxylate-O2 atoms. Thereby, a supramolecular layer is formed parallel to (100), Fig. 2a. Within this framework are benzene-1,2-diamine-N-H \cdots Cl hydrogen bonds involving all the amine-H atoms. This has the result that each chloride anion accepts four $N-H\cdots$ Cl hydrogen bonds and, to a first approximation exists in a flat, bowl-shaped environment defined by a CdH_4 'donor set'. Layers stack along the a axis with no directional interactions between them, Fig. 2b.

Given this observation, it was thought worthwhile to perform a Hirshfeld surface analysis to probe the molecular packing in more detail. The results of this analysis are discussed in the next section.

4. Hirshfeld surface analysis

The Hirshfeld surfaces calculated for (I) provide further insight into the supramolecular associations in the crystal; the calculations were performed according to a recent publication (Jotani et al., 2017). The presence of bright-red spots appearing near water-H atoms, H1W and H2W, and carboxylate oxygen atoms, O1 and O2, on the Hirshfeld surface mapped over d_{norm} in Fig. 3, result from the O-H \cdots O hydrogen bonds between these atoms, Table 2. The faint-red spots appearing near each of diamine-hydrogen atoms, H1N– H4N, and those near the Cl1 atom represent the formation of the four comparatively weak $N-H\cdots$ Cl interactions. The donors and acceptors of above intermolecular interactions can also be viewed as blue and red regions around the respective

Figure 4

A view of the Hirshfeld surface for (I) mapped over the electrostatic potential in the range -0.164 to +0.204 a.u. The red and blue regions represent negative and positive electrostatic potentials, respectively.

A view of the Hirshfeld surface for (I) mapped with the shape-index property about a reference molecule showing intermolecular $O-H\cdots O$ and $N-H\cdots$ Cl contacts as well as short interatomic $H\cdots H$ contacts as black, white and sky-blue dashed lines, respectively.

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Table 4 Summary of short inter-atomic contacts (\hat{A}) in (I) .

Contact	Distance	Symmetry operation
$H1W\cdots H3A$	2.32	$x, -1 + y, z$
$H8 \cdot H8$	2.38	$1-x, -y, 1-z$
$O1W \cdots H3A$	2.64	$x, -1 + y, z$

atoms on the Hirshfeld surface mapped over the calculated electrostatic potential in Fig. 4. The immediate environment about a reference molecule within the shape-index mapped Hirshfeld surface highlighting intermolecular $O-H \cdot \cdot O$, N- $H\cdots$ Cl interactions and short interatomic $H\cdots H$ contacts is illustrated in Fig. 5.

The overall two-dimensional fingerprint plot, Fig. 6a, and those delineated into $H \cdot \cdot H$, O $\cdot \cdot H/H \cdot \cdot \cdot C$ Cl $\cdot \cdot \cdot H/H \cdot \cdot \cdot C$ and $C \cdots H/H \cdots C$ contacts (McKinnon et al., 2007) are illustrated in Fig. 6b–e, respectively. The significant contributions from interatomic $O \cdot \cdot H/H \cdot \cdot \cdot O$ and $Cl \cdot \cdot \cdot H/H \cdot \cdot \cdot Cl$ contacts to the Hirshfeld surfaces, see data in Table 3, result from the

Figure 6

(a) The full two-dimensional fingerprint plot for (I) and fingerprint plots delineated into (b) $H \cdot \cdot H$, (c) $O \cdot \cdot H/H \cdot \cdot O$, (d) $Cl \cdot \cdot \cdot H/H \cdot \cdot \cdot Cl$ and (e) $C \cdot \cdot H/H \cdot \cdot \cdot C$ contacts.

involvement of water, diamine, chloride and carboxylate residues in the intermolecular interactions. The relatively high contribution from these atoms decreases the relative importance of interatomic $H \cdots H$ contacts, *i.e.* to 45.4%, to the Hirshfeld surface. The presence of a short interatomic $H \cdot \cdot \cdot H$ contact between water-H1W and methyl-H3A, Table 4, also has an influence upon the molecular packing as shown in Fig. 5. In the fingerprint plot delineated into $H \cdot \cdot \cdot H$ contacts, Fig. 6b, this is viewed as the distribution of points at $d_e + d_i <$ sum of their van der Waals radii, *i.e.* 2.40 Å . Another short interatomic $H \cdot \cdot \cdot H$ contact listed in Table 4, involving benzene-H8 atoms lying at the surfaces of the layers stacked along the a axis appear to have little impact upon the packing. The intermolecular $O-H\cdots O$ and $N-H\cdots Cl$ hydrogen bonding are recognized as the pair of spikes at $d_e + d_i \sim 1.8$ and 2.5 Å, respectively, together with green points within the distributions in Fig. 6c and d, respectively. The points related to short inter-atomic O \cdots H contact between water-O1W and methyl-H3A mentioned above are merged in the plot, Fig. 6c. It can be seen from the fingerprint plot delineated into $C \cdots H/H \cdots C$ contacts, Fig. 6e, that although these contacts make a significant contribution of 11.2% to the dumbbell-shaped Hirshfeld surface due to the presence of benzene-C atoms, the molecular packing results in inter-atomic $C \cdot \cdot H/H \cdot \cdot \cdot C$ separations longer than van der Waals contact distances, hence they exert a negligible effect in the crystal. The low contribution from other contacts listed in Table 3 have little effect in the structure due to their large inter-atomic separations.

5. Database survey

A search of the crystallographic literature (Groom et al., 2016) was undertaken in order to find closely related structures to (I). Reflecting the interest in these structures, there were nearly 50 examples with the adipato dianion. In each case, the dianion bridged two Cd^H cations *via* chelating interactions in all but one example. Often, the dicarboxylate ligand also bridged other Cd^H cations, *i.e.* was found to be coordinating in μ_3 - and μ_4 -modes. The most closely related structure in the literature is illustrated in Scheme 2, i.e. (II) (Che et al., 2013).

The coordination geometry for one of the independent Cd^{II} atoms in (II), being defined by two carboxylate-O atoms, derived from a tri-anionic μ_2 -benzene-1,3,5-tricarboxylato ligand, two nitrogen atoms from a chelating imidazo[4,5-f]-

Table 5 Experimental details.

Crystal data

Computer programs: CrysAlis PRO (Agilent, 2013), SHELXS97 (Sheldrick, 2008), SHELXL2014 (Sheldrick, 2015), ORTEP-3 for Windows (Farrugia, 2012), DIAMOND (Brandenburg, 2006) and publCIF (Westrip, 2010).

[1,10]phenanthroline ligand, chlorido and water-O atoms resembles that found in (I); this is illustrated on the left-hand side of Scheme 2. The difference between (I) and (II) is that in (II), the chlorido ligand is bridging, leading to a one-dimensional coordination polymer.

6. Synthesis and crystallization

Benzene-1,2-diamine (0.4324 g, 4 mmol) was slowly added to an aqueous solution (15 ml) of $CdCl₂·2H₂O$ (0.4026 g, 2 mmol) resulting in a yellow solution. The mixture was stirred for about 1 h when adipic acid (0.2923 g, 2 mmol) in MeOH (10 ml) was added. The mixture then was stirred for a further 3 h. The resultant solution was reduced and left for crystallization. Brown crystals of (I) were obtained after a few weeks and analysed directly.

7. Refinement details

Crystal data, data collection and structure refinement details are summarized in Table 5. The carbon-bound H-atoms were placed in calculated positions $(C-H = 0.95-0.99 \text{ Å})$ and were included in the refinement in the riding model approximation, with $U_{\text{iso}}(H)$ set to 1.2 $U_{\text{eq}}(C)$. The O-bound and N-bound Hatoms were located in difference-Fourier maps but were refined with distance restraints of $O-H = 0.84 \pm 0.01$ Å and $N-H = 0.88\pm0.01$ Å, and with $U_{iso}(H)$ set to $1.5U_{eq}(O)$ and $1.2U_{eq}(N)$. The maximum and minimum residual electron density peaks of 1.15 and 0.69 e \AA^{-3} , respectively, were located 0.90 and 0.87 \AA from the Cd^{II} cation.

Acknowledgements

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supporting information

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(*µ***2-Adipato-***κ***⁴** *O***,***O***′:***O***′′ ,***O***′′′)bis[aqua(benzene-1,2-diamine-**

*κ***2** *N***,***N***′)chloridocadmium]: crystal structure and Hirshfeld surface analysis**

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Computing details

Data collection: *CrysAlis PRO* (Agilent, 2013); cell refinement: *CrysAlis PRO* (Agilent, 2013); data reduction: *CrysAlis PRO* (Agilent, 2013); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL2014* (Sheldrick, 2015); molecular graphics: *ORTEP*-*3 for Windows* (Farrugia, 2012) and *DIAMOND* (Brandenburg, 2006); software used to prepare material for publication: *publCIF* (Westrip, 2010).

 $(\mu_2$ -Adipato- κ^4 O,O':O",O'")bis[aqua(benzene-1,2-diamine- $\kappa^2 N$,N')chloridocadmium]

Crystal data

 $\left[Cd_2Cl_2(C_6H_8O_4)(C_6H_8N_2)_2(H_2O)_2\right]$ $M_r = 692.14$ Monoclinic, *P*21/*c* $a = 20.4710(8)$ Å $b = 5.5578(2)$ Å $c = 10.7910(3)$ Å β = 98.122 (3)[°] $V = 1215.42(7)$ Å³ $Z = 2$

Data collection

Agilent Technologies SuperNova Dual diffractometer with Atlas detector Radiation source: SuperNova (Mo) X-ray Source Mirror monochromator Detector resolution: 10.4041 pixels mm-1 *ω* scan Absorption correction: multi-scan (CrysAlis PRO; Agilent, 2013)

Refinement

Refinement on *F*² Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.028$ $wR(F^2) = 0.065$ $S = 1.04$ 3393 reflections 163 parameters 6 restraints

 $F(000) = 684$ $D_x = 1.891$ Mg m⁻³ Mo *Kα* radiation, $\lambda = 0.71073$ Å Cell parameters from 7493 reflections θ = 3.8–30.0° μ = 2.01 mm⁻¹ $T = 100 \text{ K}$ Prism, brown $0.33 \times 0.22 \times 0.10$ mm

 $T_{\text{min}} = 0.842, T_{\text{max}} = 1.000$ 15862 measured reflections 3393 independent reflections 2992 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.039$ $\theta_{\text{max}} = 30.3^{\circ}, \theta_{\text{min}} = 3.0^{\circ}$ $h = -28 \rightarrow 26$ $k = -7 \rightarrow 7$ $l = -14 \rightarrow 14$

H atoms treated by a mixture of independent and constrained refinement $w = 1/[\sigma^2 (F_o^2) + (0.0312P)^2 + 0.8274P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\text{max}}$ < 0.001 $Δρ_{max} = 1.15 e Å⁻³$ Δ*ρ*min = −0.69 e Å−3

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

Atomic displacement parameters (Å2)

supporting information

Geometric parameters (Å, º)

Symmetry code: (i) −*x*, −*y*+2, −*z*.

Hydrogen-bond geometry (Å, º)

Symmetry codes: (ii) *x*, *y*−1, *z*; (iii) *x*, −*y*−1/2, *z*−1/2; (iv) *x*, −*y*−1/2, *z*−3/2; (v) *x*, −*y*+1/2, *z*−3/2.