

catena-Poly[[bis(μ -3-dimethylamino-*N*-salicylidenepropylamine)isothiocyanato-dicadmium(II)]- μ -isothiocyanato]**Wen-Xuan Cai, Qi-Wei Zhang,
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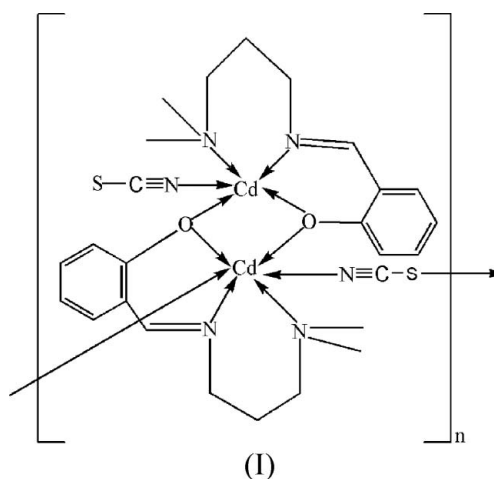
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Key indicatorsSingle-crystal X-ray study
 $T = 273$ K
Mean $\sigma(\text{C}-\text{C}) = 0.006$ Å
Disorder in main residue
 R factor = 0.028
 wR factor = 0.075
Data-to-parameter ratio = 19.4For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The deprotonated 3-dimethylamino-*N*-salicylidenepropylamine Schiff base chelates to a Cd atom in the polymeric title compound with a dinuclear building block, $[\text{Cd}_2(\text{NCS})_2(\text{C}_{12}\text{H}_{17}\text{N}_2\text{O})_2]_n$. The O atom of the Schiff base also engages in bridging. The bridging NCS group links adjacent dinuclear entities into a helical chain that runs along the b axis of the monoclinic unit cell.

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The deprotonated *N*-(salicylidene)-3-dimethylaminopropylamine Schiff base chelates to copper in the isothiocyanate complex, which exists as a centrosymmetric dinuclear compound having Cu in a square-pyramidal geometry (Lin & Feng, 2005). We present here the crystal structure of the analogous polymeric title compound with a dinuclear building block, (I), with Cd as the metal atom.



In compound (I), the deprotonated *N*-(salicylidene)-3-(dimethylamino)propylamine Schiff bases are each *O,N,N'*-chelated to a Cd atom (Fig. 1). The O atoms of the Schiff bases also engage in bridging. The set of two O and two N atoms comprises an approximate square around the Cd atom, above which lies the NCS group. For one Cd site, the geometry is octahedral, as the NCS group behaves as a bridge [$\text{Cd}-\text{N} = 2.310$ (3) and $\text{Cd}-\text{S} = 2.8165$ (8) Å], the bridging linking adjacent dinuclear entities into a helical chain that runs along the b axis of the monoclinic unit cell (Fig. 2). For the other Cd site, the geometry is square pyramidal, with the N atom of the NCS group at the apical site. However, the geometry is distorted towards octahedral, owing to the proximity of the S atom that is involved in bridging [$\text{Cd}-\text{N} = 2.219$ (3) and $\text{Cd}\cdots\text{S} = 3.2814$ (10) Å].

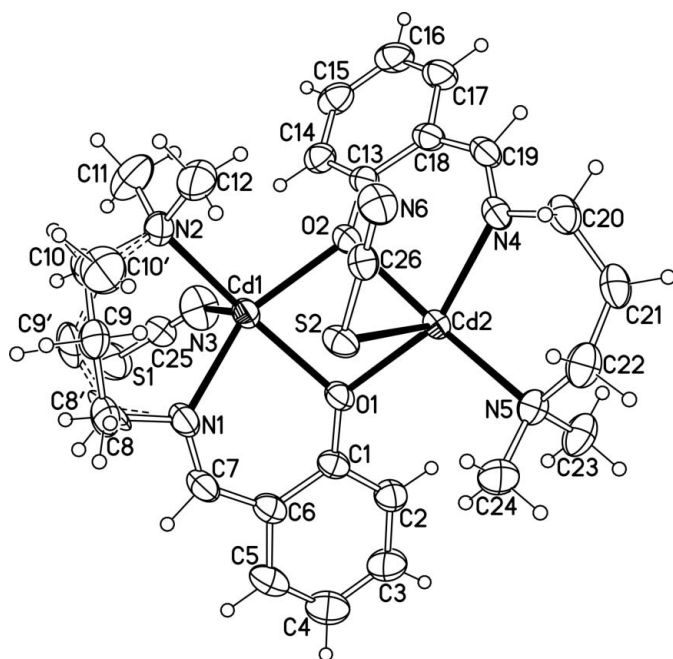


Figure 1

The asymmetric unit of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii. The minor component of the disordered trimethylene linkage is shown with dashed bonds. [Symmetry code (i): $\frac{3}{2} - x, y - \frac{1}{2}, \frac{3}{2} - z$.]

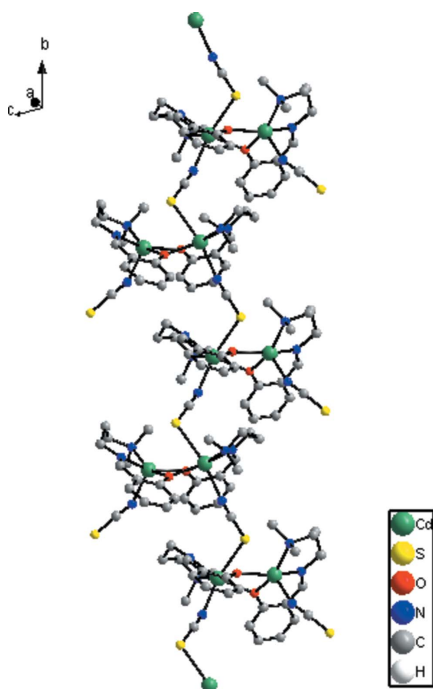


Figure 2

The helical chain structure of (I). H atoms have been omitted

Experimental

The Schiff base ligand *N*-(salicylidene)-3-dimethylaminopropylamine was prepared by refluxing 3-dimethylamino-1-propylamine (0.125 ml, 1.0 mmol) and salicylaldehyde (0.105 ml, 1.0 mmol) in ethanol (10 ml)

for 30 min. $\text{Cd}(\text{CH}_3\text{COO})_2 \cdot 2\text{H}_2\text{O}$ (0.267 g, 1.0 mmol) was then added dropwise. A yellow mixture was obtained by refluxing for about an hour, and this was cooled to room temperature and filtered. Yellow crystals of (I) suitable for X-ray diffraction were obtained by slow evaporation of the yellow filtrate in air (yield 47%).

Crystal data

$[\text{Cd}_2(\text{NCS})_2(\text{C}_{12}\text{H}_{17}\text{N}_2\text{O})_2]$
 $M_r = 751.55$
 Monoclinic, $P2_1/c$
 $a = 14.9782$ (15) Å
 $b = 12.7398$ (13) Å
 $c = 21.7317$ (15) Å
 $\beta = 132.148$ (4)°
 $V = 3074.5$ (5) Å³

$Z = 4$
 $D_x = 1.624$ Mg m⁻³
 Mo $K\alpha$ radiation
 $\mu = 1.55$ mm⁻¹
 Prism, yellow
 $0.40 \times 0.38 \times 0.20$ mm

Data collection

Bruker APEX-II area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.543, T_{\max} = 0.733$

20224 measured reflections
 7274 independent reflections
 5081 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.021$
 $\theta_{\max} = 27.9^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.028$
 $wR(F^2) = 0.075$
 $S = 1.06$
 7274 reflections
 374 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0345P)^2 + 0.5071P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.95$ e Å⁻³
 $\Delta\rho_{\min} = -0.78$ e Å⁻³

Table 1

Selected bond lengths (Å).

Cd1—N3	2.219 (3)	Cd2—O1	2.2586 (17)
Cd1—O2	2.2420 (18)	Cd2—O2	2.2759 (19)
Cd1—O1	2.2446 (19)	Cd2—N4	2.287 (2)
Cd1—N1	2.286 (2)	Cd2—N6 ⁱ	2.310 (3)
Cd1—N2	2.344 (2)	Cd2—N5	2.352 (2)
Cd1...Cd2	3.4302 (4)	Cd2—S2	2.8165 (8)

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

H atoms were positioned geometrically ($\text{C}-\text{H} = 0.93\text{--}0.97$ Å), and were included in the refinement in the riding-model approximation, with $U_{\text{iso}}(\text{H})$ restrained to 1.2–1.5 times $U_{\text{eq}}(\text{C})$. The methyl groups were rotated to fit the electron density. One of the trimethylene linkages is disordered over two positions and the site occupancy factors refined to 0.61 (1):0.39 (1). The pairs of bond distances (N1–C8/N1–C8', C8–C9/C8'–C9', C9–C10/C9'–C10' and C10–N2/C10'–N2) were restrained to within 0.01 Å of each other.

Data collection: *SMART* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2004); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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