metal-organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 292 KMean $\sigma(\text{C}-\text{C}) = 0.004 \text{ Å}$ R factor = 0.048 wR factor = 0.126 Data-to-parameter ratio = 14.8

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

catena-Poly[[[4-(1*H*-1,3,7,8-tetraazacyclopenta-[*I*]phenanthren-2-yl)phenol]manganese(II)]*µ*-benzene-1,4-dicarboxylato]

In the title compound, $[Mn(C_8H_4O_4)(C_{19}H_{12}N_4O)_2]_n$ or $[Mn(1,4-BDC)(L)_2]_n$, where 1,4-BDC is the benzene-1,4-dicarboxylate dianion and L is 4-(1*H*-1,3,7,8-tetraazacyclopenta[*l*]phenanthren-2-yl)phenol, the Mn^{II} atom is six-coordinated by two bidentate L and two monodentate 1,4-BDC ligands, resulting in a distorted *trans*-MnO₂N₄ octahedral geometry. The Mn^{II} atoms are bridged by the 1,4-BDC ligands, forming a single-chain polymer structure. In addition, neighboring single chains interact through π - π interactions, generating a two-dimensional network. Mn has site symmetry $\overline{1}$; the complete 1,4-BDC unit is also built up by inversion symmetry.

Comment

Coordination polymers with a variety of supramolecular structures are attracting increasing interest because of their novel topologies and potential applications as functional materials (Eddaoudi *et al.*, 2001). The 1,10-phenanthroline (phen) ligand has been widely used to build novel supramolecular architectures through its aromatic π - π interactions (Chen & Liu, 2002). As one important derivative of phen, 4-(1*H*-1,3,7,8-tetraazacyclopenta[*I*]phenanthren-2-yl)-phenol (*L*), was recently shown to form a coordination polymer with a distinctive supramolecular architecture (Che *et al.*, 2006). We selected benzene-1,4-dicarboxylic acid (1,4-H₂BDC) to act as a metal-metal linker in its deprotonated form and *L* as a secondary ligand, generating the title compound, [Mn(1,4-BDC)(*L*)₂], (I), a new coordination polymer, which is reported here.



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In compound (I), the Mn^{II} atom is coordinated by four N atoms from two bidentate L ligands and two O atoms from

Received 26 May 2006

Accepted 26 May 2006



Figure 1

The asymmetric unit of compound (I), together with symmetryequivalent atoms to complete the Mn coordination and the 1,4-BDC ligand, showing the atomic numbering scheme. Displacement ellipsoids are drawn at the 30% probability level (arbitrary spheres for the H atoms). [Symmetry codes: (i) -x, -y, 1 - z; (ii) -x, -y, -z.]

two 1,4-BDC ligands in a distorted cis-MnO₂N₄ octahedral geometry (Fig. 1). Mn1 has site symmetry $\overline{1}$ and the complete 1,4-BDC dianion is also generated by inversion. The Mn-O and mean Mn-N distances [2.296 (2) Å] in (I) (Table 1) are comparable to those observed for [Mn(1,4-BDC) $(L')_2$ [L' is (2-phenyl-1H-1,3,7,8-tetraazacyclopenta[l]phenanthrene] (Che, 2006). Neighbouring Mn^{II} atoms are bridged by 1,4-BDC ligands, forming a single-chain structure (Fig. 2). The C-O bond lengths (Table 1) indicate that the carboxylate bonding is essentially localized.

Neighbouring single chains in (I) are connected through π - π interactions between L ligands with a π - π stacking distance of 3.53 Å, resulting in a two-dimensional supramolecular structure (Fig. 3) (Noveron et al., 2002). Finally, interchain N- $H \cdots O$ and $O - H \cdots O$ hydrogen bonds (Table 2) complete the structure of (I).

Experimental

Ligand L was synthesized according to the literature method of Steck & Day (1943). An ethanolic solution (8 ml) of L (0.5 mmol) was added slowly to an aqueous solution (6 ml) of MnCl₂·2H₂O (0.5 mmol) and 1,4-H₂BDC (1 mmol) with stirring at refluxing temperature. The resulting solution was filtered and the filtrate was allowed to stand in air at room temperature for two weeks, yielding vellow crystals of (I) (49% yield based on Mn).



Figure 2 View of the single-chain structure of (I). H atoms have been omitted.





Crystal data

| | IZ 000 0 (4) Å3 |
|---------------------------------------|---|
| $[Mn(C_8H_4O_4)(C_{19}H_{12}N_4O)_2]$ | V = 909.0 (4) A ² |
| $M_r = 843.70$ | Z = 1 |
| Triclinic, $P\overline{1}$ | $D_x = 1.541 \text{ Mg m}^{-3}$ |
| $a = 9.1936 (18) \text{\AA}$ | Mo $K\alpha$ radiation |
| b = 10.515 (2) Å | $\mu = 0.43 \text{ mm}^{-1}$ |
| c = 11.056 (2) Å | T = 292 (2) K |
| $\alpha = 61.77 \ (3)^{\circ}$ | Block, yellow |
| $\beta = 78.55 \ (3)^{\circ}$ | $0.32 \times 0.30 \times 0.27 \text{ mm}$ |
| $\gamma = 75.90 \ (3)^{\circ}$ | |
| | |

Data collection

| Rigaku R-AXIS RAPID | 8954 measured reflections |
|--|--|
| diffractometer | 4106 independent reflections |
| ω scans | 3064 reflections with $I > 2\sigma(I)$ |
| Absorption correction: multi-scan | $R_{\rm int} = 0.032$ |
| (ABSCOR; Higashi, 1995) | $\theta_{\rm max} = 27.5^{\circ}$ |
| $T_{\min} = 0.865, \ T_{\max} = 0.891$ | |
| | |

| $w = 1/[\sigma^2(F_0^2) + (0.0636P)^2]$ |
|--|
| + 0.1901P] |
| where $P = (F_0^2 + 2F_c^2)/3$ |
| $(\Delta/\sigma)_{\rm max} = 0.001$ |
| $\Delta \rho_{\rm max} = 1.51 \text{ e } \text{\AA}^{-3}$ |
| $\Delta \rho_{\rm min} = -0.36 \text{ e } \text{\AA}^{-3}$ |
| |
| |

Table 1

Refinement

Selected geometric parameters (Å, °).

| 1.224 (3) | Mn1-N1 | 2.283 (2) |
|-------------|--|--|
| 1.283 (3) | Mn1-N2 | 2.309 (2) |
| 2.1635 (16) | | |
| 73.22 (8) | | |
| | 1.224 (3) 1.283 (3) 2.1635 (16) 73.22 (8) | 1.224 (3) Mn1–N1 1.283 (3) Mn1–N2 2.1635 (16) 73.22 (8) |

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Table 2

Hydrogen-bond geometry (Å, $^{\circ}$).

| $D - H \cdots A$ | <i>D</i> -H | $H \cdots A$ | $D \cdots A$ | $D - H \cdots A$ |
|-------------------------|-------------|---------------------|--------------|------------------|
| N4-H4···O3 ⁱ | 0.86 | 1.92 | 2.775 (3) | 173 |
| $O1-H1A\cdots O2^n$ | 0.82 | 1.93 | 2.654 (2) | 147 |
| Summature and an (i) | | (3) $n + 1$ $n + 1$ | - 1 | |

Symmetry codes: (i) -x, -y + 1, -z; (ii) x + 1, y + 1, z - 1.

All H atoms were positioned geometrically (N-H = 0.86 Å and C-H = 0.93 Å) and refined as riding, with $U_{iso}(H) = 1.2U_{eq}(\text{carrier})$. The highest residual density peak is located 0.90 Å from atom H18.

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *PROCESS-AUTO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL-Plus* (Sheldrick, 1990); software used to prepare material for publication: *SHELXL97*. The authors thank Jilin Normal University for supporting this work.

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