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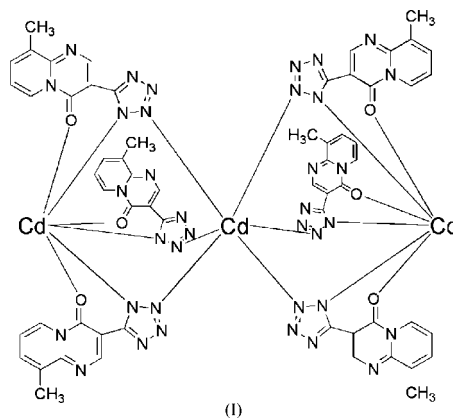
Key indicators

Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.004$ Å
 R factor = 0.032
 wR factor = 0.079
Data-to-parameter ratio = 15.4For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Hexakis[μ -9-methyl-3-(1*H*-tetrazol-5-*io*)-
4*H*-pyrido[1,2-*a*]pyrimidin-4-onato(2-)]-
tricadmium(II)The centrosymmetric molecule of the title Cd^{II} complex, $[\text{Cd}_3(\text{C}_{10}\text{H}_7\text{N}_6\text{O})_6]$, is located on a threefold axis. The pyridopyrimidine ligands bridge neighbouring Cd atoms, forming a trinuclear complex. π - π Stacking is observed between neighbouring complex molecules.

Received 20 December 2005

Accepted 12 January 2006

Comment

Recent developments in polynuclear cadmium complexes concern their interesting topologies and properties (Zheng *et al.*, 2003; Bu *et al.*, 2005; Chandrasekhar *et al.*, 2005; Liu, 2005; Yong *et al.*, 2005). We present here the structure of the title trinuclear Cd^{II} complex incorporating a tetrazole ligand, $[\text{Cd}_3(\text{pemirolast})_6]$ [pemirolast = 9-methyl-3-(1*H*-tetrazol-5-*io*)-4*H*-pyrido[1,2-*a*]pyrimidin-4-one, which is an anti-allergic drug], (I).

The molecular structure of (I) is shown in Fig. 1. The trinuclear molecule has a threefold axis of symmetry with three Cd atoms located on the threefold axis. The central Cd1 atom is also located on an inversion centre and coordinated by six N atoms from six pemirolast ligands, with the Cd1–N bond length of 2.335 (2) Å; the bond angles around the Cd1 centre are close to 90°, showing an almost ideal octahedral geometry. The terminal Cd2 atom is coordinated by three N atoms and three O atoms from three pemirolast ligands, with the Cd2–N and Cd2–O bond lengths of 2.294 (2) and 2.351 (2) Å, respectively. Atom Cd2 has a slightly distorted octahedral coordination geometry (Table 1). In the trinuclear units, all of the pemirolast ligands adopt an *O,N,N*-tridentate chelating-bridging coordination mode, using the tetrazole groups to bridge adjacent Cd atoms, giving rise to the trinuclear complex.

The C2-containing pyridopyrimidine is nearly parallel with the C2¹-containing one [symmetry code: (i) $2 - x, 1 - x + y, \frac{1}{2} -$

z], the dihedral angle being $4.89(11)^\circ$ (Fig. 2). The distances of atoms on the C2-pyridoperimidine from the mean plane of the C2ⁱ-pyridoperimidine are 3.294(3) (C5), 3.409(3) (C6), 3.392(3) (C8) and 3.272(3) Å (N6). These clearly suggest the existence of π - π stacking in the crystal structure of (I).

Experimental

A mixture of Cd(ClO₄)₂·6H₂O (0.05 mmol) and the potassium salt of 9-methyl-3-(1*H*-tetrazol-5-yl)-4*H*-pyrido[1,2-*a*]pyrimidin-4-one (0.10 mmol) was placed in a heavy-walled Pyrex tube containing pyridine (0.05 ml), ethanol (0.30 ml) and H₂O (0.10 ml). The tube was frozen in liquid N₂, sealed under vacuum, and then heated at 393 K for 2 d. **Caution:** Cd(ClO₄)₂·6H₂O is potentially explosive and should be used with care. Colourless crystals suitable for X-ray diffraction analysis were collected, washed with ethanol, and dried in air.

Crystal data

[Cd₃(C₁₀H₇N₆O)₆]
M_r = 1700.53
 Trigonal, *R*3̄*c*
a = 18.302(3) Å
c = 31.099(6) Å
V = 9021(3) Å³
Z = 6
D_x = 1.878 Mg m⁻³

Mo *K*α radiation
 Cell parameters from 15120 reflections
 $\theta = 1.8$ – 28.3°
 $\mu = 1.14$ mm⁻¹
T = 293(2) K
 Block, colourless
 0.44 × 0.39 × 0.30 mm

Data collection

Siemens SMART CCD diffractometer
 φ and ω scans
 Absorption correction: none
 15306 measured reflections
 2466 independent reflections

1874 reflections with $I > 2\sigma(I)$
*R*_{int} = 0.043
 $\theta_{\max} = 28.3^\circ$
h = -24 → 18
k = -10 → 24
l = -40 → 40

Refinement

Refinement on *F*²
R [*F*² > 2σ(*F*²)] = 0.032
wR [*F*²] = 0.079
S = 0.99
 2466 reflections
 160 parameters

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0385P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.002$
 $\Delta\rho_{\max} = 0.88$ e Å⁻³
 $\Delta\rho_{\min} = -0.66$ e Å⁻³

Table 1
 Selected bond lengths (Å).

Cd1—N2	2.335(2)	Cd2—O1	2.3514(19)
Cd2—N1	2.294(2)		

Aromatic H atoms were placed in calculated positions, with C—H = 0.93 Å, and refined in the riding mode, with *U*_{iso}(H) = 1.2*U*_{eq}(C). The methyl H atoms were placed in calculated positions, with C—H = 0.96 Å, the torsion angles refined, and *U*_{iso}(H) = 1.5*U*_{eq}(C).

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1998); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine

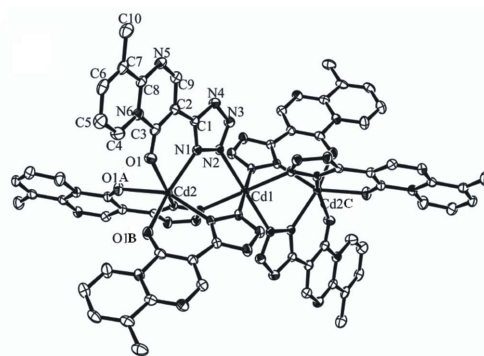


Figure 1
 The molecular structure of (I) with 50% probability displacement ellipsoids [symmetry codes: (A) $-y, x - y, z$; (B) $-x + y, -x, z$; (C) $-x, -y, -z$]. H atoms have been omitted for clarity.

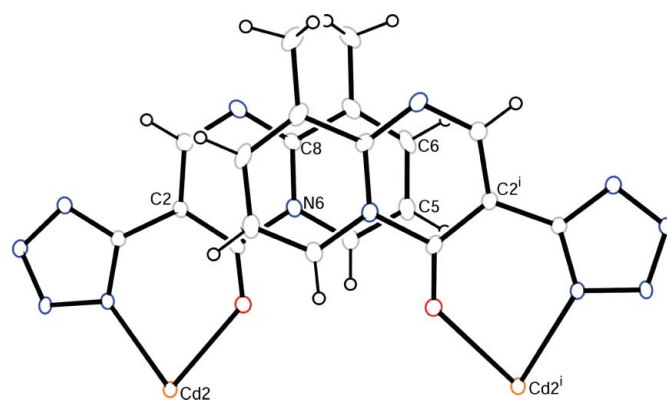


Figure 2
 A diagram showing the π - π stacking [symmetry code: (i) $2 - x, 1 - x + y, \frac{1}{2} - z$].

structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1998); software used to prepare material for publication: SHELXTL.

We are grateful for financial support from NSFC (No. 20472078).

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