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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.004 Å R factor = 0.044 wR factor = 0.109 Data-to-parameter ratio = 17.9

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. In the title mononuclear zinc(II) compound, $[Zn(C_{16}H_{36}N_4)]$ -(ClO₄)₂, the Zn²⁺ atom lies on an inversion center and is coordinated by four N atoms from the macrocyclic ligand. The coordination geometry is slightly distorted square planar.

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Comment

In preparing $Zn(ClO_4)_2$ complexes with ethylenediamine in acetone, we isolated some crystals of the title compound, (I), unexpectedly. However, we failed to repeat the experiment. The complex can also be obtained according to a published method (Zhu *et al.*, 1999).



In the complex cation, the Zn^{II} atom, on an inversion center, is coordinated by four N atoms of the macrocyclic ligand (Fig. 1). The metal atom and the four N atoms are coplanar. The Zn-N bond lengths are close to those in similar zinc(II) complexes (Zhu *et al.*, 2004; You *et al.*, 2004). The perchlorate anions are not coordinated.

Experimental

The ligand *meso*-5,5,7,12,12,14-hexamethyl-1,4,8,11-tetraazacyclotetradecane was prepared according to the literature method of Hay *et al.* (1975). Equimolar amounts of the ligand and $Zn(ClO_4)_2 \cdot 9H_2O$ were mixed in water in a flask and the solution was kept in air for one week. Large colorless prismatic crystals were precipitated after about 80% of the solvent had evaporated. They were filtered off, washed with water three times and dried in air (yield 24%).

Crystal data [Zn(C16H32N4)](ClO4)2 $D_{\rm r} = 1.622 {\rm Mg m}^{-3}$ $M_r = 544.74$ Mo $K\alpha$ radiation Monoclinic, $P2_1/c$ Cell parameters from 1989 a = 10.323 (2) Å reflections b = 10.630 (2) Å = 4.8-27.5° $\mu = 1.38 \text{ mm}^{-1}$ c = 11.018 (2) Å $\beta = 111.71 \ (3)^{\circ}$ T = 293 (2) K V = 1123.3 (4) Å² Prism, colorless $0.45 \times 0.35 \times 0.35 \text{ mm}$ Z = 2

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Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{min} = 0.580, T_{max} = 0.625$ 9499 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.044$ $wR(F^2) = 0.109$ S = 1.142645 reflections 148 parameters H atoms treated by a mixture of independent and constrained refinement 2645 independent reflections 2448 reflections with $I > 2\sigma(I)$ $R_{int} = 0.024$ $\theta_{max} = 28.2^{\circ}$ $h = -13 \rightarrow 12$ $k = -13 \rightarrow 13$ $l = -14 \rightarrow 14$

 $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0422P)^{2} + 1.108P]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3 (\Delta/\sigma)_{max} = 0.029 \Delta\rho_{max} = 0.44 \text{ e} \text{ Å}^{-3} \Delta\rho_{min} = -0.45 \text{ e} \text{ Å}^{-3}$

All the H atoms attached to C atoms were positioned geometrically and constrained to ride on their parent atoms, with C–H distances of 0.96 or 0.97 Å and with $U_{iso}(H) = 1.2U_{eq}(C)$ or $1.5U_{eq}(C)$. The H atom attached to N1 was refined isotropically.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997*a*); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997*a*); molecular graphics: *SHELXTL* (Sheldrick, 1997*b*); software used to prepare material for publication: *SHELXTL*.

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Figure 1

The molecular structure of (I), showing 30% probability displacement ellipsoids and the atom-numbering scheme. Unlabeled atoms are related to labeled atoms by (2 - x, 2 - y, -z).

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