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Key indicators

Single-crystal X-ray study
 T = 298 K
 Mean $\sigma(\text{C}-\text{C}) = 0.007 \text{ \AA}$
 Disorder in solvent or counterion
 R factor = 0.068
 wR factor = 0.185
 Data-to-parameter ratio = 12.4

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

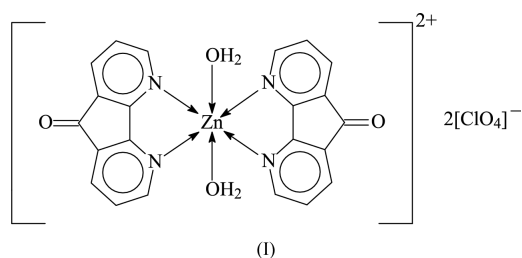
Diaquabis(4,5-diazafluoren-9-one- $\kappa^2\text{N,N}'$)-zinc(II) diperchlorate

The Zn atom in the title compound, $[\text{Zn}(\text{C}_{11}\text{H}_6\text{N}_2\text{O})_2(\text{H}_2\text{O})_2](\text{ClO}_4)_2$, lies on a center of symmetry in the crystal structure, and it is chelated by the heterocyclic ligands [$\text{Zn}-\text{O} = 2.151(4)$ and $2.276(4) \text{ \AA}$]. The coordinated water molecule interacts with the perchlorate ions to form a hydrogen-bonded layer structure. The compound is isomorphous with the Cu analog, whose structure has been detailed by Menon & Rajasekharan [*Polyhedron* (1998), **17**, 2463–2476].

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Comment

Although 4,5-diazafluoren-9-one sometimes does not exhibit its chelating behavior, it is known to bind copper(II) if the anionic group is the perchlorate group. The crystal structure of diaquabis(4,5-diazafluoren-9-one)copper diperchlorate has been described in detail (Gu *et al.*, 2002; Menon & Rajasekharan, 1998; Zhao *et al.*, 2003; Zhang, Zhao *et al.*, 2003). The present centrosymmetric Zn analog, (I) (Fig. 1), is isomorphous with the Cu compound. With chloride as the counterion, the ligand does not behave as a chelate; the ligand is instead protonated in $(\text{C}_6\text{H}_{12}\text{N}_2\text{O})_2(\text{ZnCl}_4) \cdot 2\text{H}_2\text{O}$ (Zhang, Liu *et al.*, 2003).



Experimental

One drop of tris(2-aminoethyl)amine and three drops of perchloric acid were added to an ethanol solution of zinc dichloride (0.14 g, 1 mmol). 4,5-Diazafluoren-9-one (0.18 g, 1 mmol) dissolved in ethanol was added, and the mixture heated for 1 h. Yellow crystals of (I) separated from the solution after several days.

Crystal data

$[\text{Zn}(\text{C}_{11}\text{H}_6\text{N}_2\text{O})_2(\text{H}_2\text{O})_2](\text{ClO}_4)_2$	$D_x = 1.774 \text{ Mg m}^{-3}$
$M_r = 664.66$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 6381 reflections
$a = 8.1493(4) \text{ \AA}$	$\theta = 2.3\text{--}28.1^\circ$
$b = 11.5615(5) \text{ \AA}$	$\mu = 1.28 \text{ mm}^{-1}$
$c = 13.2055(6) \text{ \AA}$	$T = 298(2) \text{ K}$
$\beta = 90.662(1)^\circ$	Irregular block, yellow
$V = 1244.1(1) \text{ \AA}^3$	$0.31 \times 0.27 \times 0.25 \text{ mm}$
$Z = 2$	

Data collection

Bruker SMART APEX area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.440$, $T_{\max} = 0.724$
 14 093 measured reflections

2848 independent reflections
 2431 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.040$
 $\theta_{\max} = 27.5^\circ$
 $h = -10 \rightarrow 10$
 $k = -15 \rightarrow 14$
 $l = -17 \rightarrow 17$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.068$
 $wR(F^2) = 0.185$
 $S = 1.10$
 2848 reflections
 230 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.1043P)^2 + 0.991P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.60 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.44 \text{ e } \text{\AA}^{-3}$

Table 1

Selected geometric parameters (\AA , $^\circ$).

Zn1—O1w	2.073 (3)	Zn1—N2	2.151 (4)
Zn1—N1	2.276 (4)		
O1w—Zn1—O1w ⁱ	180.0	N1—Zn1—N1 ⁱ	180.0
O1w—Zn1—N1	88.9 (2)	N1—Zn1—N2	80.8 (1)
O1w—Zn1—N1 ⁱ	91.1 (2)	N1—Zn1—N2 ⁱ	99.2 (1)
O1w—Zn1—N2	90.6 (1)	N2—Zn1—N2 ⁱ	180.0
O1w—Zn1—N2 ⁱ	89.4 (1)		

Symmetry code: (i) $1 - x, 1 - y, 1 - z$.

Table 2

Hydrogen-bonding geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1w—H1w1 \cdots O2	0.84 (1)	1.96 (4)	2.667 (7)	140 (6)
O1w—H1w2 \cdots O4 ⁱⁱⁱ	0.84 (1)	2.03 (2)	2.80 (1)	152 (4)
O1w—H1w2 \cdots O4 ⁱⁱⁱ	0.84 (1)	1.84 (4)	2.54 (1)	138 (6)

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

The aromatic H atoms were positioned geometrically [$C-H = 0.93 \text{ \AA}$ and $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C)$] and were allowed to ride on their parent atoms in the riding-model approximation. The water H atoms were located and refined with distance restraints $O-H = 0.85 (1)$ and $H\cdots H = 1.39 (1) \text{ \AA}$. The perchlorate ion is disordered over two sites sharing a common Cl atom, and it was refined as two ions, with distance restraints $Cl-O = 1.44 (1)$ and $O\cdots O = 2.35 (1) \text{ \AA}$. The displacement parameters of the O atoms were restrained to be approximately isotropic. The perchlorate group refined to a ratio of $0.62 (1)/0.38 (1)$.

Data collection: SMART (Bruker, 2001); cell refinement: SMART; data reduction: SAINT (Bruker, 2001); program(s) used to solve

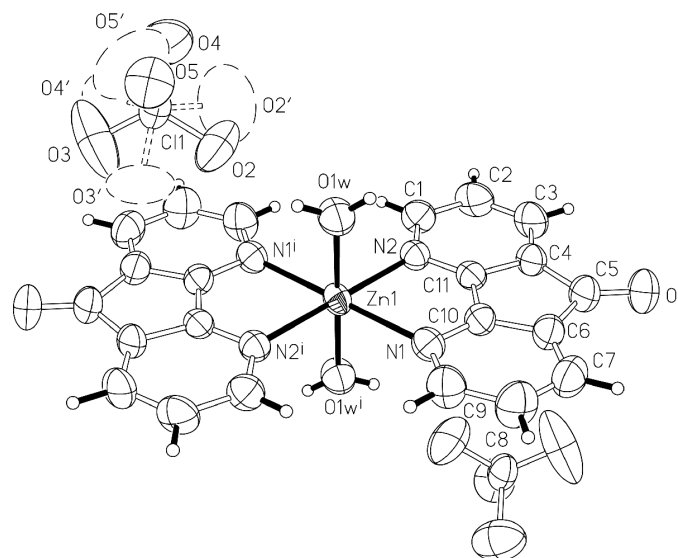


Figure 1

ORTEPII (Johnson, 1976) plot of (I); displacement ellipsoids are drawn at the 50% probability level. H atoms are drawn as spheres of arbitrary radii. Both disorder components of the perchlorate anion are shown. [Symmetry code: (i) $1 - x, 1 - y, 1 - z$.]

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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