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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=293 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.004 \AA$
H -atom completeness $76 \%$
Disorder in solvent or counterion
$R$ factor $=0.055$
$w R$ factor $=0.186$
Data-to-parameter ratio $=16.0$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## A magnesium(II) complex of 1,10-phenanthroline-2,9dicarboxylate

The title compound, triaqua(1,10-phenanthroline-2,9-dicarboxylato)magnesium(II) dihydrate, $\left[\mathrm{Mg}(\mathrm{PDA})\left(\mathrm{H}_{2} \mathrm{O}\right)_{3}\right] \cdot 2 \mathrm{H}_{2} \mathrm{O}$, $\left(\mathrm{H}_{2} \mathrm{PDA}\right.$ is 1,10-phenanthroline-2,9-dicarboxylic acid, $\mathrm{C}_{14} \mathrm{H}_{8^{-}}$ $\mathrm{N}_{2} \mathrm{O}_{4}$ ) has twofold crystallographic symmetry. The Mg atom is in a distorted pentagonal bipyramidal coordination environment with two N atoms and two O atoms from PDA and one O atom from a water molecule forming the pentagonal plane, and two O atoms from two water molecules occupying axial positions. The crystal structure comprises an infinite twodimensional network of hydrogen-bonded molecules.

## Comment

1,10-Phenanthroline-2,9-dicarboxylic acid ( $\mathrm{H}_{2} \mathrm{PDA}$ ) has been used as a simple sensitizing species of luminescent lanthanide ion chelates for analytical applications in aqueous solution (Sammes \& Yahioglu, 1994; Mullins et al., 1996). However, investigations of $\mathrm{H}_{2} \mathrm{PDA}$ complexes with metal ions, such as $\mathrm{Fe}^{\mathrm{II}}$ and $\mathrm{Eu}^{\mathrm{II}}$, have been limited to spectroscopic characterizations in aqueous solution (König \& Ritter, 1981; Templeton \& Pollak, 1989; Sammes et al., 1992; Dyson et al., 1999). To our knowledge, no examples of $\mathrm{Mg}^{\mathrm{II}}$ complexes of $\mathrm{H}_{2}$ PDA have been characterized in the solid state. We have prepared the $\mathrm{Mg}^{\text {II }}$ complex of $\mathrm{H}_{2} \mathrm{PDA}$, (I), and report its crystal structure here.

(I)

The title compound (Fig. 1) is located on a twofold axis of symmetry which passes through the Mg and O 3 atoms. The seven-coordinated Mg atom is in a distorted pentagonal bipyramidal geometry. Two N and two O atoms from PDA and one O atom from a water molecule define the pentagonal plane, and the two axial positions are occupied by O atoms derived from two water molecules.

Important bond distances and angles are presented in Table 1. The bond distances between Mg and the PDA donor atoms $\left[\begin{array}{lll}\mathrm{Mg}-\mathrm{O} 1 & 2.3080(17) ~ \AA \\ \mathrm{~A}\end{array} \mathrm{and} \mathrm{Mg}-\mathrm{N} 1 \quad 2.2994\right.$ (19) $\AA$ ] are significantly longer than those to coordinated water mol-

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Figure 1
The structure of the title compound with the atom-numbering scheme. The displacement ellipsoids are drawn at the $50 \%$ probability level (Johnson, 1976). The non-coordinated water molecule has been omitted for clarity. [Symmetry code: (i) $-x+\frac{3}{4},-y+\frac{3}{4}, z$ ].


Figure 2
Packing diagram of (I) viewed along [100]. Hydrogen bonds are indicated by dashed lines. Displacement ellipsoids are shown at the $30 \%$ probability level. All H atoms and the disordered water molecules with lower site occupancy have been omitted for clarity.
ecules [ $\mathrm{Mg}-\mathrm{O} 32.055$ (2) $\AA$ and $\mathrm{Mg}-\mathrm{O} 42.0777$ (18) $\AA$ ]. This is probably due to the high rigidity of PDA as well as the high affinity of the $\mathrm{Mg}^{\text {II }}$ ion to water molecules. The carboxylate groups of the PDA ligand are almost coplanar with the phenanthroline unit as indicated by the $\mathrm{O} 1-\mathrm{C} 1-\mathrm{C} 2-\mathrm{N} 1$ torsion angle of 1.9 (3) ${ }^{\circ}$.

The complexes are inter-connected by hydrogen bonds between the coordinated water molecules, O3 and O4, and the
carboxylate O atoms of adjacent PDA with interaction distances of 2.774 (2) and 2.745 (3) A (Table 2); the equatorial water molecule is hydrogen bonded with the coordinated carboxylate O atoms and the axial water molecules interact with the carbonyl O atoms.

As illustrated in Fig. 2, the complexes associate along the $a$ axis and form columns in the crystal structure. Non-coordinated water molecules also participate in hydrogen bonds and serve to connect the complex units along the $b$ axis. Stacking interactions between centrosymmetrically related phenanthroline units are observed with a plane-to-plane separation of 3.360 (4) Å.

## Experimental

$\mathrm{H}_{2}$ PDA was synthesized according to the literature (König \& Ritter, 1981). The title compound was crystallized by slow evaporation from the methanol solution prepared by the reaction of equimolar amounts of $\mathrm{H}_{2} \mathrm{PDA}$ and $\mathrm{MgSO}_{4}$.

## Crystal data

$\left[\mathrm{Mg}\left(\mathrm{C}_{14} \mathrm{H}_{6} \mathrm{~N}_{2} \mathrm{O}_{4}\right)\left(\mathrm{H}_{2} \mathrm{O}\right)_{3}\right] \cdot 2 \mathrm{H}_{2} \mathrm{O} \quad$ Mo $K \alpha$ radiation
$M_{r}=380.60$
Orthorhombic, Fidd
$a=7.4194$ (12) £
$b=19.044$ (3) $\AA$
$c=46.943(7) \AA$
$V=6632.8(18) \AA^{3}$
$V=6632.8(18) \AA^{3}$
$Z=16$
$D_{x}=1.525 \mathrm{Mg} \mathrm{m}^{-3}$

## Data collection

| CCD area detector diffractometer | $R_{\text {int }}=0.066$ |
| :--- | :--- |
| $\varphi$ and $\omega$ scans | $\theta_{\max }=28.3^{\circ}$ |
| 10406 measured reflections | $h=-9 \rightarrow 9$ |
| 2063 independent reflections | $k=-25 \rightarrow 25$ |
| 1385 reflections with $I>2 \sigma(I)$ | $l=-62 \rightarrow 42$ |

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.055$
$w R\left(F^{2}\right)=0.186$
$S=1.06$
2063 reflections
129 parameters
H atoms treated by a mixture of independent and constrained refinement

Table 1
Selected geometric parameters $\left(\AA,{ }^{\circ}\right)$.

| $\mathrm{Mg}-\mathrm{O} 3$ | $2.055(2)$ | $\mathrm{Mg}-\mathrm{N} 1$ | $2.2994(19)$ |
| :--- | :---: | :--- | :--- |
| $\mathrm{Mg}-\mathrm{O} 4$ | $2.0777(18)$ | $\mathrm{Mg}-\mathrm{O} 1$ | $2.3080(17)$ |
|  |  |  |  |
| $\mathrm{O} 3-\mathrm{Mg}-\mathrm{O} 4$ | $89.81(5)$ | $\mathrm{O} 3-\mathrm{Mg}-\mathrm{O} 1$ | $77.65(5)$ |
| $\mathrm{O} 4^{\mathrm{i}}-\mathrm{Mg}-\mathrm{O} 4$ | $179.63(11)$ | $\mathrm{O} 4-\mathrm{Mg}-\mathrm{O} 1$ | $87.41(7)$ |
| $\mathrm{O} 4-\mathrm{Mg}-\mathrm{N} 1$ | $89.28(7)$ | $\mathrm{N} 1-\mathrm{Mg}-\mathrm{O} 1$ | $68.04(6)$ |
| $\mathrm{N} 1-\mathrm{Mg}-\mathrm{N} 1^{\mathrm{i}}$ | $68.64(10)$ |  |  |

Symmetry codes: (i) $\frac{3}{4}-x, \frac{3}{4}-y, z$.

Table 2
Hydrogen-bonding geometry $\left(\AA,{ }^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots \mathrm{A}$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :---: | :---: | :---: | :---: | :---: |
| $\mathrm{O} 3-\mathrm{H} 3 A \cdots \mathrm{O} 1^{\text {i }}$ | 0.991 (1) | 1.791 (2) | 2.774 (2) | 170.6 (1) |
| $\mathrm{O} 4-\mathrm{H} 4 A \cdots \mathrm{O} 2^{\text {ii }}$ | 0.856 (2) | 1.891 (2) | 2.745 (3) | 175.8 (1) |
| $\mathrm{O} 4-\mathrm{H} 4 \mathrm{~B} \cdots \mathrm{O} 5$ | 0.922 (2) | 1.960 (7) | 2.863 (6) | 165.7 (4) |
| $\mathrm{O} 4-\mathrm{H} 4 \mathrm{~B} \cdots \mathrm{O}^{\prime}$ | 0.922 (2) | 1.89 (1) | 2.80 (1) | 169 (2) |
| $\mathrm{O} 5 \cdots \mathrm{O} 2^{\text {iii }}$ |  |  | 2.866 (7) |  |
| $\mathrm{O} 5^{\prime} \cdots \mathrm{O} 2^{\text {iii }}$ |  |  | 2.71 (2) |  |
| O5 $\cdots \mathrm{OS}^{\text {iv }}$ |  |  | 2.84 (1) |  |
| O5 ${ }^{\prime} \cdots \mathrm{OF}^{\text {,iv }}$ |  |  | 2.73 (3) |  |

Symmetry codes: (i) $x-\frac{1}{2}, \frac{3}{4}-y, \frac{1}{4}-z$; (ii) $x-1, y, z$; (iii) $\frac{5}{4}-x, y, \frac{1}{4}-z$; (iv) $x, \frac{5}{4}-y, \frac{1}{4}-z$.

The $\mathrm{C}-\mathrm{H}$ atoms were added at their calculated positions $\left[U_{\text {iso }}=\right.$ $1.2 U_{\text {eq }}(\mathrm{C})$ ] and refined using a riding model. The H atoms of the coordinated water molecules were located from a difference map but were not refined. The non-coordinated water molecule, O5, is disordered over two sites with occupancies of 0.7 for O 5 and 0.3 for $\mathrm{O}^{\prime}$; H atoms were not included for this molecule.

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

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