

Qian Yang, Shan Gao\* and  
Li-Hua HuoCollege of Chemistry and Chemical Technology,  
Heilongjiang University, Harbin 150080,  
People's Republic of ChinaCorrespondence e-mail:  
shangao67@yahoo.com

## Key indicators

Single-crystal X-ray study  
 $T = 296$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å  
 $R$  factor = 0.046  
 $wR$  factor = 0.122  
Data-to-parameter ratio = 13.8For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.Hexaaquamagnesium(II) benzene-1,4-dioxy-  
acetate

The title complex, alternatively called hexaaquamagnesium(II) *p*-phenylenebis(oxyacetate),  $[\text{Mg}(\text{H}_2\text{O})_6](\text{C}_{10}\text{H}_8\text{O}_6)$ , has been synthesized by the reaction of  $\text{Mg}(\text{NO}_3)_2 \cdot 6\text{H}_2\text{O}$  and benzene-1,4-dioxyacetic acid in an aqueous solution. The  $\text{Mg}^{\text{II}}$  atom is coordinated by six water molecules to form an octahedron [ $\text{Mg}-\text{O} = 2.0433(16)-2.1281(18)$  Å]. The cations and anions, which lie on centres of symmetry, are linked by hydrogen bonds into a three-dimensional network structure.

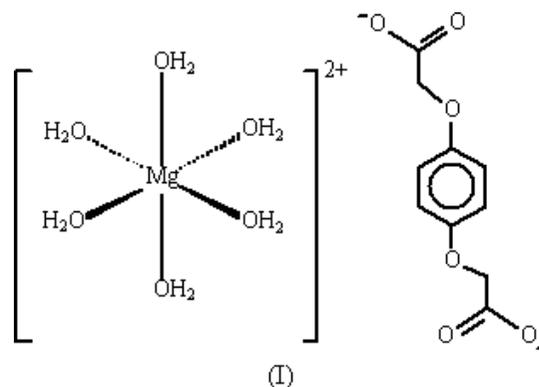
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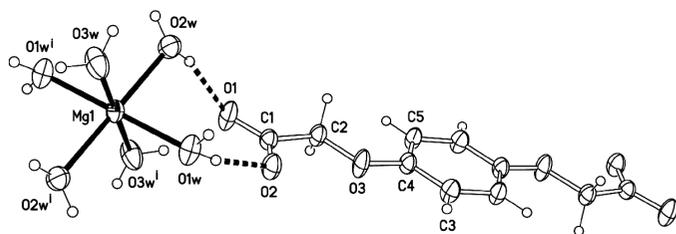
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## Comment

Benzene-1,4-dioxyacetic acid is known as a good candidate for the construction of supramolecular complexes; it can form regular hydrogen bonds by functioning as either hydrogen-bond donor or acceptor (Gao *et al.*, 2004). We have previously reported the structures of hexaaquacobalt(II) benzene-1,4-dioxyacetate (Liu, Huo *et al.*, 2004) and hexaaquamanganese(II) benzene-1,4-dioxyacetate (Liu, Gao *et al.*, 2004). Recently, we found that benzene-1,4-dioxyacetic acid forms a hexaaquamagnesium(II) benzene-1,4-dioxyacetate complex, (I), under similar reaction conditions and that this is isomorphous with the  $\text{Co}^{\text{II}}$  and  $\text{Mn}^{\text{II}}$  analogues. Previous structural descriptions of the  $\text{Co}^{\text{II}}$  and  $\text{Mn}^{\text{II}}$  complexes apply to the present isostructural complex, (I) (Fig. 1).



The  $\text{Mg}^{\text{II}}$  atom is six-coordinate in an octahedral environment. The anion is almost planar (r.m.s. deviation = 0.01 Å); the oxyacetate group and aromatic ring are essentially planar, with a torsion angle of  $-179.1(15)^\circ$  ( $\text{C}4-\text{O}3-\text{C}2-\text{C}1$ ). The cation and anion both lie on inversion centres and are linked by extensive hydrogen bonds into a three-dimensional supramolecular network structure (Table 2).



**Figure 1**  
ORTEP plot (Johnson, 1976) of the title complex, with displacement ellipsoids drawn at the 30% probability level and hydrogen bonds shown as dashed lines. Symmetry codes are as in Table 1.

**Experimental**

Benzene-1,4-dioxyacetic acid was prepared by the nucleophilic reaction of chloroacetate and hydroquinone under basic conditions, following the method described for the synthesis of benzene-1,2-dioxyacetic acid (Mirci, 1988). The title complex was prepared by the addition of  $Mg(NO_3)_2 \cdot 6H_2O$  (5.13 g, 20 mmol) and KSCN (0.97 g, 10 mol) to an aqueous solution (25 ml) of benzene-1,4-dioxyacetic acid (4.52 g, 20 mmol). The resulting solution was stirred and the pH was adjusted to 7 with 0.1 M NaOH solution. Colourless crystals were separated from the filtered solution after several days. Analysis calculated for  $C_{10}H_{20}MgO_{12}$ : C 33.69, H 5.65%; found: C 33.80, H 5.61%.

*Crystal data*

$[Mg(H_2O)_6](C_{10}H_8O_6)$   $Z = 1$   
 $M_r = 356.57$   $D_x = 1.557 \text{ Mg m}^{-3}$   
 Triclinic,  $P\bar{1}$  Mo  $K\alpha$  radiation  
 Cell parameters from 3689 reflections  
 $a = 5.5677(11) \text{ \AA}$   
 $b = 6.3662(13) \text{ \AA}$   
 $c = 11.620(2) \text{ \AA}$   
 $\alpha = 102.11(3)^\circ$   
 $\beta = 95.59(3)^\circ$   
 $\gamma = 106.68(3)^\circ$   
 $V = 380.26(16) \text{ \AA}^3$   
 $\theta = 3.5\text{--}27.5^\circ$   
 $\mu = 0.18 \text{ mm}^{-1}$   
 $T = 296(2) \text{ K}$   
 Prism, colourless  
 $0.37 \times 0.25 \times 0.18 \text{ mm}$

*Data collection*

Rigaku R-AXIS RAPID diffractometer 1710 independent reflections  
 1321 reflections with  $I > 2\sigma(I)$   
 $\omega$  scans  $R_{int} = 0.027$   
 Absorption correction: multi-scan (*ABSCOR*; Higashi, 1995)  $\theta_{max} = 27.5^\circ$   
 $T_{min} = 0.947, T_{max} = 0.968$   $h = -7 \rightarrow 7$   
 3747 measured reflections  $k = -8 \rightarrow 8$   
 $l = -14 \rightarrow 14$

*Refinement*

Refinement on  $F^2$   $w = 1/[\sigma^2(F_o^2) + (0.0638P)^2 + 0.1773P]$   
 $R[F^2 > 2\sigma(F^2)] = 0.046$  where  $P = (F_o^2 + 2F_c^2)/3$   
 $wR(F^2) = 0.122$   $(\Delta/\sigma)_{max} < 0.001$   
 $S = 1.02$   $\Delta\rho_{max} = 0.49 \text{ e \AA}^{-3}$   
 1710 reflections  $\Delta\rho_{min} = -0.30 \text{ e \AA}^{-3}$   
 124 parameters  
 H atoms treated by a mixture of independent and constrained refinement

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

Mg1—O1w	2.0433 (16)	O1—C1	1.237 (2)
Mg1—O2w	2.0677 (16)	O2—C1	1.265 (3)
Mg1—O3w	2.1281 (18)		
O1w—Mg1—O1w <sup>i</sup>	180	O2w <sup>i</sup> —Mg1—O2w	180
O1w—Mg1—O2w	92.50 (7)	O2w <sup>i</sup> —Mg1—O3w	93.09 (7)
O1w—Mg1—O3w	92.03 (7)	O2w—Mg1—O3w	86.91 (7)
O1w <sup>i</sup> —Mg1—O2w	87.50 (7)	O3w <sup>i</sup> —Mg1—O3w	180
O1w <sup>i</sup> —Mg1—O3w	87.97 (7)		

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

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

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O1w—H1w1 $\cdots$ O2 <sup>ii</sup>	0.85 (2)	2.03 (2)	2.823 (2)	155 (3)
O1w—H1w2 $\cdots$ O1 <sup>i</sup>	0.84 (2)	1.90 (2)	2.722 (2)	163 (3)
O2w—H2w1 $\cdots$ O3w <sup>ii</sup>	0.86 (2)	2.16 (2)	2.993 (2)	164 (2)
O2w—H2w2 $\cdots$ O2	0.86 (2)	1.89 (2)	2.740 (2)	168 (3)
O3w—H3w1 $\cdots$ O1	0.84 (2)	2.06 (2)	2.869 (2)	161 (2)
O3w—H3w2 $\cdots$ O2 <sup>iii</sup>	0.85 (2)	1.87 (2)	2.707 (2)	170 (2)
O3w—H3w2 $\cdots$ O3 <sup>iii</sup>	0.85 (2)	2.58 (2)	3.094 (3)	120 (2)

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

Water H atoms were located in a difference map and refined with O—H distance restraints of 0.85 (1)  $\text{\AA}$  and with  $U_{iso}(H) = 1.5U_{eq}(O)$ . The other H atoms were placed in calculated positions [ $C-H = 0.93$  (aromatic) or 0.97  $\text{\AA}$  (aliphatic) and  $U_{iso}(H) = 1.2U_{eq}(C)$ ] and were refined using the riding-model approximation.

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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