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#### **Key indicators**

Single-crystal X-ray study T = 298 KMean  $\sigma$ (C–C) = 0.004 Å R factor = 0.029 wR factor = 0.079 Data-to-parameter ratio = 11.4

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

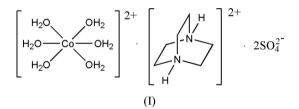
# 1,4-Diazoniabicyclo[2.2.2]octane hexaaquacobalt(II) bis(sulfate)

The title compound,  $(C_6H_{14}N_2)[Co(H_2O)_6](SO_4)_2$ , consists of hexaaquacobalt(II) cations, 1,4-diazabicyclo[2,2,2]octane cations and sulfate anions, linked by  $O-H\cdots O$  hydrogen bonds. The geometry around the  $Co^{2+}$  cation is octahedral.

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### Comment

In the synthesis of crystal structures by design, the assembly of molecular units in predefined arrangements is a key goal (Desiraju, 1995, 1997; Braga *et al.*, 1998). Directional intermolecular interactions are the primary tools in achieving this goal and hydrogen bonding is currently the best among these (Zaworotko, 1997; Braga & Grepioni, 2000). In this paper, we report the structure of the title compound, (I).



The asymmetric unit of (I) consists of a 1,4-diazabicyclo-[2,2,2]octane cation, a hexaaquacobalt(II) cation and two sulfate anions. The Co<sup>2+</sup> cation lies in a general position and the geometry around it is octahedral, with bonds to six water molecules (Fig. 1 and Table 1). The coordinated water molecules, 1,4-diazabicyclo[2,2,2]octane cations and sulfate anions interact through hydrogen bonds (Table 2), generating a three-dimensional network (Fig. 2).

#### **Experimental**

Cobalt(II) sulfate heptahydrate (0.06 g, 0.2 mmol) was dissolved in water (10 ml) and the solution was mixed with a dimethylformamide solution (10 ml) of 5-sulfoisophthalic acid monosodium salt (0.11 g, 0.4 mmol) and 1,4-diazabicyclo[2,2,2]octane (0.05 g, 0.4 mmol). The reaction mixture was filtered and allowed to stand. Pink prism-shaped crystals of (I) separated from the solution after about three months. As shown by the present crystal structure analysis, no component of the 5-sulfoisophthalic acid monosodium salt was incorporated into the product.

## Crystal data

(C<sub>6</sub>H<sub>14</sub>N<sub>2</sub>)[Co(H<sub>2</sub>O)<sub>6</sub>](SO<sub>4</sub>)<sub>2</sub>  $D_x = 1.820 \text{ Mg m}^{-3}$ Mo  $K\alpha$  radiation  $M_r = 473.34$ Monoclinic,  $P2_1/c$ Cell parameters from 3079 reflections a = 12.0488 (8) Å b = 12.1842 (8) Å = 1.7-25.1°  $\mu = 1.31 \text{ mm}^{-1}$ c = 12.1390 (8) Å  $\beta = 104.20 \ (1)^{\circ}$ T = 298 (2) K V = 1727.8 (2) Å Prism, pink  $0.29 \times 0.26 \times 0.20 \text{ mm}$ Z = 4

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# metal-organic papers

#### Data collection

Bruker APEX area-detector
diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan
(SADABS; Bruker, 2002)
$T_{\min} = 0.69, \ T_{\max} = 0.77$
8951 measured reflections

#### Refinement

Refinement on $F^2$
$R[F^2 > 2\sigma(F^2)] = 0.029$
$wR(F^2) = 0.079$
S = 1.06
3079 reflections
269 parameters
H atoms treated by a mixture of
independent and constrained
refinement

3079 independent reflections 2920 reflections with  $I > 2\sigma(I)$  $R_{int} = 0.020$  $\theta_{max} = 25.1^{\circ}$  $h = -14 \rightarrow 12$  $k = -14 \rightarrow 14$  $l = -14 \rightarrow 11$ 

$$\begin{split} &w = 1/[\sigma^2(F_o^2) + (0.0428P)^2 \\ &+ 1.1451P] \\ &where \ P = (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{max} = 0.001 \\ \Delta\rho_{max} = 0.29 \ e^{-3} \\ \Delta\rho_{min} = -0.44 \ e^{-3} \\ Extinction correction: SHELXTL \\ (Bruker, 2002) \\ Extinction coefficient: 0.0642 (18) \end{split}$$

### Table 1

Selected geometric parameters (Å, °).

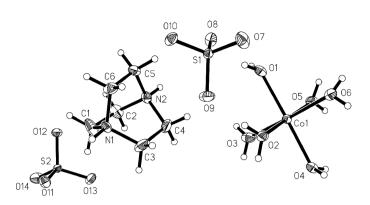
Co1-O2	2.0705 (17)	Co1-O1	2.0816 (16)
Co1-O4	2.0728 (16)	Co1-O3	2.1053 (17)
Co1-O5	2.0798 (17)	Co1-O6	2.1466 (16)
O2-Co1-O4	86.86 (7)	O5-Co1-O3	88.36 (8)
O2-Co1-O5	175.82 (7)	O1-Co1-O3	89.01 (7)
O4-Co1-O5	93.86 (7)	O2-Co1-O6	89.04 (7)
O2-Co1-O1	91.75 (7)	O4-Co1-O6	92.45 (7)
O4-Co1-O1	178.43 (7)	O5-Co1-O6	86.82 (7)
O5-Co1-O1	87.57 (7)	O1-Co1-O6	88.27 (7)
O2-Co1-O3	95.75 (8)	O3-Co1-O6	174.56 (7)
O4-Co1-O3	90.39 (7)		

Table 2			
Hydrogen-bond	geometry	(Å,	°).

	• • • •			
$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O6−H6C···O12 <sup>i</sup>	0.83 (3)	1.95 (3)	2.780 (2)	177 (4)
O2−H2C···O9	0.82 (3)	1.94 (3)	2.752 (2)	171 (4)
$O1 - H1C \cdot \cdot \cdot O8$	0.81 (3)	1.98 (3)	2.776 (2)	168 (4)
$O2-H2D\cdots O13^{ii}$	0.79 (3)	1.98 (3)	2.753 (2)	164 (4)
$O4-H4D\cdots O13^{i}$	0.84 (3)	1.85 (3)	2.691 (2)	174 (4)
$O1 - H1D \cdots O10^{iii}$	0.80 (3)	1.90 (3)	2.691 (2)	169 (4)
$O5-H5C\cdots O12^{iv}$	0.79 (3)	1.98 (3)	2.765 (2)	171 (4)
$O6-H6D\cdots O8^{v}$	0.83 (3)	2.16 (3)	2.985 (2)	173 (4)
$O3-H3C\cdots O9^{iii}$	0.82 (3)	1.98 (3)	2.802 (3)	179 (4)
$O5-H5D\cdots O10^{v}$	0.81(2)	1.89 (3)	2.688 (2)	169 (4)
$O4-H4C\cdots O11^{iv}$	0.82 (3)	1.90 (3)	2.698 (2)	167 (4)
O3−H3D···O14 <sup>vi</sup>	0.81(2)	2.01 (3)	2.790 (2)	160 (4)
$N2-H2N\cdots O7^{iii}$	0.86 (2)	1.86 (3)	2.685 (3)	159 (3)
$N1 - H1N \cdot \cdot \cdot O14^{vii}$	0.89 (2)	2.03 (3)	2.796 (3)	145 (3)
$N1 - H1N \cdot \cdot \cdot O12^{vii}$	0.89 (2)	2.23 (3)	2.995 (3)	145 (3)

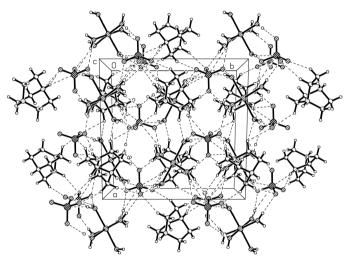
Symmetry codes: (i)  $x, -y + \frac{3}{2}, z + \frac{1}{2}$ ; (ii)  $-x + 1, y + \frac{1}{2}, -z + \frac{3}{2}$ ; (iii)  $x, -y + \frac{3}{2}, z - \frac{1}{2}$ ; (iv) x, y + 1, z; (v)  $-x, y + \frac{1}{2}, -z + \frac{3}{2}$ ; (vi) -x + 1, -y + 1, -z + 1; (vii)  $x, -y + \frac{1}{2}, z + \frac{1}{2}$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

Water and amine H atoms were located in a difference Fourier map and refined isotropically, with O–H and N–H distance restraints of 0.82 (3) and 0.86 (3) Å, respectively. All other H atoms were positioned geometrically and allowed to ride on their parent atoms at distances of 0.97 Å, with  $U_{iso}(H) = 1.2U_{eq}$ (parent atom).



#### Figure 1

The asymmetric unit of (I), showing the atom-numbering scheme and with displacement ellipsoids at the 50% probability level.



#### Figure 2

A perspective view of the three-dimensional network of (I), assembled *via* hydrogen bonds, which are shown as dashed lines.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Bruker, 2002); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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