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## Key indicators

Single-crystal X-ray study
$T=293 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.041$
$w R$ factor $=0.081$
Data-to-parameter ratio $=13.7$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## trans-Diaquatetrakis(4,4'-methylenediphenyl-amine- $\kappa N$ )cobalt(II) dinitrate dihydrate

The hydrothermally prepared title compound, $\left[\mathrm{Co}\left(\mathrm{C}_{13} \mathrm{H}_{14}{ }^{-}\right.\right.$ $\left.\left.\mathrm{N}_{2}\right)_{4}\left(\mathrm{H}_{2} \mathrm{O}\right)_{2}\right]\left(\mathrm{NO}_{3}\right)_{2} \cdot 2 \mathrm{H}_{2} \mathrm{O}$, consists of mononuclear cationic complexes of six-coordinate cobalt(II), together with nitrate counter-ions and uncoordinated water molecules. Four monodentate $4,4^{\prime}$-methylenediphenylamine (dapm) ligands and two water molecules comprise the octahedral coordination of the $\mathrm{Co}^{\mathrm{II}}$ ion, which lies on an inversion center. The compound is isostructural with the previously reported Cd and Ni analogs.

## Comment

The structure of the title compound, (I), consists of a pseudooctahedral $\mathrm{Co}^{2+}$ complex with four monodentate dapm ligands [dapm is $4,4^{\prime}$-methylenediphenylamine] coordinated through one amine N and two water molecules, as shown in Fig. 1, with two nitrate ions and two uncoordinated water molecules. The $\mathrm{Co}^{2+}$ ion lies on a crystallographic inversion center. Hydrothermally prepared (I) is isostructural with the Cd and Ni analogs obtained from room temperature evaporation of water-alcohol mixtures of the $M\left(\mathrm{NO}_{3}\right)_{2}$ salt $(M=\mathrm{Cd}, \mathrm{Ni})$ and dapm (Wang et al., 2001; Zhang et al., 2001).

(I)

## Experimental

The title compound was prepared by hydrothermal reaction of 4,4'methylenedianiline (dapm) $\quad(0.0240 \mathrm{~g}, \quad 0.12 \mathrm{mmol}) \quad$ with $\mathrm{Co}\left(\mathrm{NO}_{3}\right)_{2} \cdot 6 \mathrm{H}_{2} \mathrm{O}(0.0386 \mathrm{~g}, 0.11 \mathrm{mmol})$ in water $(0.80 \mathrm{ml})$ in an evacuated sealed Pyrex tube. The tube was heated to 358 K at 1 K


Figure 1
The structure of (I). Displacement ellipsoids are drawn at the $30 \%$ probability level. Symmetry code: (') $1-x 1,-y, 1-z$. The symmetryrelated nitrate ion and uncoordinated water molecule are not shown.

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$\min ^{-1}$, and held at that temperature for 8 h before slowly cooling ( $0.2 \mathrm{~K} \mathrm{~min}^{-1}$ ) to 303 K . Pale-orange block-shaped crystals were formed along with some pink powder. Analysis calculated for $\mathrm{C}_{52} \mathrm{H}_{64} \mathrm{CoN}_{10} \mathrm{O}_{10}$ : C 59.59 , H 6.16, N $13.36 \%$; found: C 59.25, H 5.87, N $13.22 \%$.

## Crystal data

$\left[\mathrm{Co}\left(\mathrm{C}_{13} \mathrm{H}_{14} \mathrm{~N}_{2}\right)_{4}\left(\mathrm{H}_{2} \mathrm{O}\right)_{2}\right]-$

$$
\left(\mathrm{NO}_{3}\right)_{2} \cdot 2 \mathrm{H}_{2} \mathrm{O}
$$

$$
M_{r}=1048.06
$$

$$
\begin{aligned}
& Z=1 \\
& D_{x}=1.370 \mathrm{Mg} \mathrm{~m}^{-3} \\
& \text { Mo } K \alpha \text { radiation } \\
& \text { Cell parameters from } 6696 \\
& \quad \text { reflections } \\
& \theta=2.3-26.3^{\circ} \\
& \mu=0.41 \mathrm{~mm}^{-1} \\
& T=293(2) \mathrm{K} \\
& \text { Block, pale orange } \\
& 0.42 \times 0.24 \times 0.18 \mathrm{~mm}
\end{aligned}
$$

Triclinic, $P \overline{1}$
$a=9.2265$ (5) $\AA$ 。
$b=11.6587$ (6) $\AA$
$c=11.9719$ (7) $\AA$
$\alpha=80.920(1)^{\circ}$
$\beta=87.804(1)^{\circ}$
$\gamma=89.488(1)^{\circ}$
$V=1270.73(12) \AA^{3}$
Data collection

| Bruker SMART APEX CCD | 5200 independent reflections |
| :--- | :--- |
| diffractometer | 3875 reflections with $I>2 \sigma(I)$ |
| $\omega$ and $\varphi$ scans | $R_{\text {int }}=0.030$ |
| Absorption correction: multi-scan | $\theta_{\max }=26.4^{\circ}$ |
| $(S A D A B S ;$ Bruker, 1999$)$ | $h=-11 \rightarrow 11$ |
| $T_{\min }=0.619, T_{\text {max }}=0.928$ | $k=-14 \rightarrow 14$ |
| 11803 measured reflections | $l=-14 \rightarrow 14$ |

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.041$
$w R\left(F^{2}\right)=0.081$
H atoms treated by a mixture of independent and constrained refinement
$S=0.99$
5200 reflections
379 parameters

Table 2
Hydrogen-bonding geometry ( $\AA \mathrm{A}^{\circ}$ ).

| $D-\mathrm{H} \cdots \mathrm{A}$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots \mathrm{A}$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :---: | :---: | :---: | :---: | :---: |
| $\mathrm{N} 1-\mathrm{H} 1 \mathrm{~A} \cdots \mathrm{O}^{\text {i }}$ | 0.90 (2) | 2.22 (2) | 3.114 (2) | 173.2 (18) |
| $\mathrm{N} 3-\mathrm{H} 3 \mathrm{~A} \cdots \mathrm{O} 3^{\text {ii }}$ | 0.82 (2) | 2.52 (2) | 3.263 (3) | 150.0 (19) |
| $\mathrm{O} 1-\mathrm{H} 1 W A \cdots \mathrm{O}^{\text {ii }}$ | 0.79 (2) | 1.99 (3) | 2.779 (2) | 175 (3) |
| $\mathrm{N} 4-\mathrm{H} 4 \mathrm{~B} \cdots \mathrm{O} 5^{\text {iii }}$ | 0.93 (3) | 2.29 (3) | 3.107 (3) | 147 (2) |
| $\mathrm{O} 2-\mathrm{H} 2 \mathrm{WA} \cdots \mathrm{O} 5^{\text {iii }}$ | 0.76 (4) | 2.56 (4) | 3.193 (4) | 142 (4) |
| $\mathrm{N} 2-\mathrm{H} 2 \mathrm{~B} \cdots \mathrm{O}^{\text {iv }}$ | 0.86 (3) | 2.46 (3) | 3.258 (4) | 156 (2) |
| $\mathrm{N} 4-\mathrm{H} 44 \cdots \mathrm{O} 4^{\text {v }}$ | 0.96 (2) | 2.14 (3) | 3.048 (3) | 157 (2) |
| $\mathrm{O} 1-\mathrm{H} 1 W B \cdots \mathrm{O} 2^{\text {vi }}$ | 0.83 (2) | 1.87 (3) | 2.694 (3) | 170 (2) |
| $\mathrm{O} 2-\mathrm{H} 2 W B \cdots \mathrm{~N} 2^{\mathrm{i}}$ | 0.72 (4) | 2.26 (4) | 2.958 (5) | 165 (5) |

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

H atoms attached to C atoms were geometrically idealized, with $\mathrm{C}-\mathrm{H}=0.93 \AA$ and $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C}) . \mathrm{H}$ atoms on N atoms and water O were located and refined freely with isotropic displacement parameters.

Data collection: SMART-NT (Bruker, 1999); cell refinement: SAINT-Plus-NT (Bruker, 1999); data reduction: SAINT-Plus-NT ; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: XP in SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: SHELXTL.

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