

# Bis[*trans*-dibromo(2,2-dimethylpropane-1,3-diamine- $\kappa^2N,N'$ )-chromium(III)] dibromide hydrogen perchlorate hexahydrate

Jong-Ha Choi,<sup>a</sup> Takayoshi Suzuki<sup>b</sup> and Sumio Kaizaki<sup>b\*</sup>

<sup>a</sup>Department of Chemistry, Andong National University, Andong 760-749, South Korea, and <sup>b</sup>Department of Chemistry, Graduate School of Science, Osaka University, Toyonaka 560-0043, Japan

Correspondence e-mail: kaizaki@chem.sci.osaka-u.ac.jp

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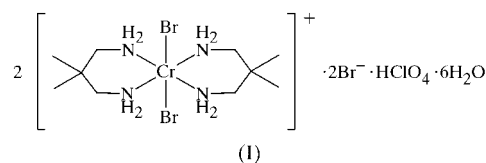
In the title compound,  $[\text{CrBr}_2(\text{C}_5\text{H}_{14}\text{N}_2)_2]_2\text{Br}_2\cdot\text{HClO}_4\cdot 6\text{H}_2\text{O}$ , there are two independent  $\text{Cr}^{\text{III}}$  complex cations which are conformational isomers of each other. The Cr atoms lie respectively on a center of symmetry and on a mirror plane and have octahedral environments, coordinated by the N atoms of two 2,2-dimethylpropane-1,3-diamine ligands and by two Br atoms in *trans* positions. The Cr–N and Cr–Br bond lengths are in the ranges 2.078 (3)–2.089 (3) and 2.4495 (9)–2.5017 (9) Å, respectively. The crystal structure consists of two  $\text{Cr}^{\text{III}}$  complex cations, two  $\text{Br}^-$  anions, a  $(\text{ClO}_4)^-$  anion and an  $[\text{H}_{13}\text{O}_6]^+$  hydrogen-bonded cluster cation.

## Comment

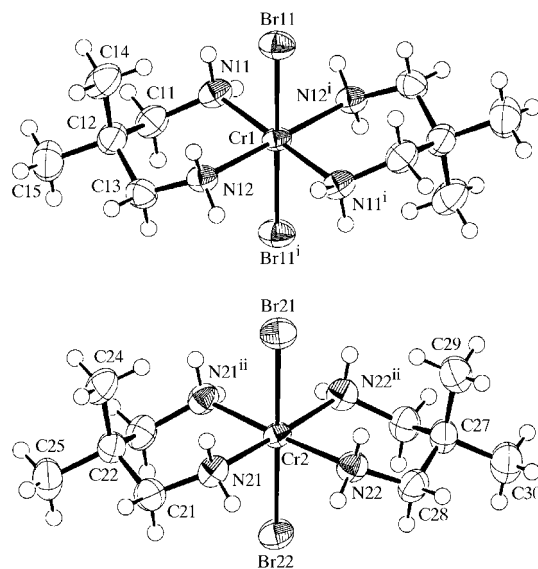
2,2-Dimethylpropane-1,3-diamine ( $\text{Me}_2\text{tn}$ ) can coordinate to a central metal ion as a bidentate ligand, forming a six-membered chelate ring, similar to the situation for unsubstituted propane-1,3-diamine (tn) (House, 1986). In addition to the conformational isomerism resulting from the six-membered chelate rings, the  $[\text{CrBr}_2(\text{tn}/\text{Me}_2\text{tn})_2]^+$  complex cations can exist as *cis* or *trans* isomers. Although electronic absorption and IR spectra have been used to identify the geometrical structures of chromium(III) complexes (Choi *et al.*, 2002; Poon & Pun, 1980), assignments based on spectral data are not always conclusive (Stearns & Armstrong, 1992). Furthermore, since reports on the structural chemistry of complexes of the type  $[\text{Cr}^{\text{III}}\text{L}_4\text{Br}_2]\text{X}$  are rare (Lisgarten *et al.*, 1990), we prepared the title complex, (I), which contains the  $[\text{CrBr}_2(\text{Me}_2\text{tn})_2]^+$  cation, and determined its crystal structure by X-ray analysis.

The structure analysis of (I) reveals the crystal to contain two crystallographically independent  $\text{Cr}^{\text{III}}$  complex moieties. As atom Cr1 is located at a crystallographic center of symmetry, this  $\text{Cr}^{\text{III}}$  complex cation has molecular  $C_i$  symmetry, while the other complex cation, containing atom

Cr2, has molecular  $C_s$  symmetry; atoms Cr2, Br21, Br22, C22, C24, C25, C27, C29 and C30 are located on a crystallographic mirror plane. It was found that in both complex cations, the two Br atoms are coordinated to the central Cr atom in a *trans* arrangement. The Cr–Br bond lengths [2.4495 (9)–2.5017 (9) Å] are close to the value found in *trans*- $[\text{CrBr}_2(\text{cyclam})]\text{Br}$  (cyclam is 1,4,8,11-tetraazacyclotetradecane; Lisgarten *et al.*, 1990). The Cr–N bond lengths in (I) are in the range 2.078 (3)–2.089 (3) Å, which are typical values for  $\text{Cr}^{\text{III}}$ –N(primary amine) bonds. All of the six-membered chelate rings formed by the  $\text{Me}_2\text{tn}$  ligand adopt chair conformations and the two complex cations are so-called conformational isomers ( $C_i$  and  $C_s$ ) due to the relative conformation of the two  $\text{Me}_2\text{tn}$  chelate rings, as shown in Fig. 1.

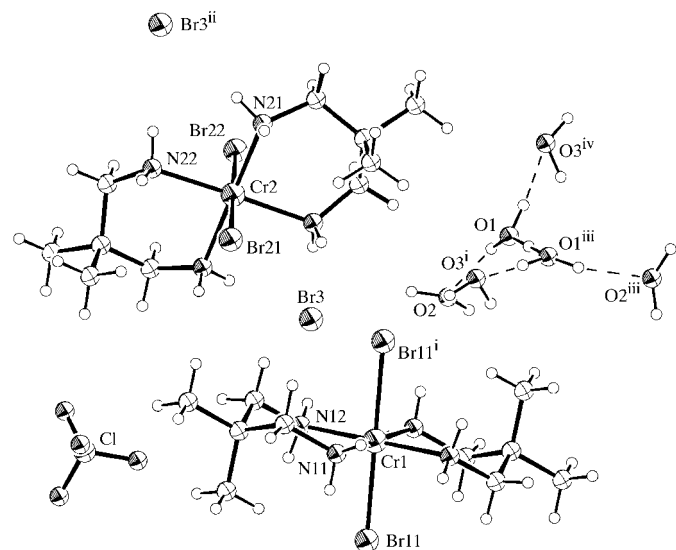


As well as the complex cation, the asymmetric unit of (I) also contains a  $\text{Br}^-$  ion (Br3), half a  $(\text{ClO}_4)^-$  anion and three O atoms (O1, O2 and O3) corresponding to the hydrated water molecules. For charge balance, a half equivalent amount of  $\text{H}^+$  per  $\text{Cr}^{\text{III}}$  complex cation should be included in the crystal. We suspected that such an  $\text{H}^+$  ion was trapped between atoms O1 and O1<sup>iii</sup> [symmetry code: (iii)  $1 - x, y, -z$ ], because the O1...O1<sup>iii</sup> distance was only 2.347 (9) Å, pointing to the existence of a strong hydrogen-bonding interaction between these atoms (Minkwitz *et al.*, 1998). Since atom O1 is also involved in hydrogen bonds with both O2 and O3<sup>iv</sup> [O1...O2 = 2.662 (7) Å and O1...O3<sup>iv</sup> = 2.667 (7) Å; symmetry code: (iv)  $\frac{1}{2} + x, \frac{1}{2} - y, z$ ], the hydrated  $\text{H}^+$  cation in



**Figure 1**  
Perspective views of the two crystallographically independent complex cations in (I). Displacement ellipsoids are drawn at the 50% probability level. [Symmetry codes: (i)  $\frac{1}{2} - x, \frac{1}{2} - y, -z$ ; (ii)  $x, -y, z$ .]

the crystal of (I) forms an  $[\text{H}_{13}\text{O}_6]^+$  cluster cation (Fig. 2). Although both Eigen-type and chain-like structures of  $[\text{H}_9\text{O}_4]^+$  clusters are often observed (Frank & Reiss, 1997; Hassaballa *et al.*, 1998), the H-shaped  $[\text{H}_{13}\text{O}_6]^+$  cluster cation found in the crystal of (I) has, to the best of our knowledge, not been reported previously.



**Figure 2**  
A view of a complete set of ions in (I). [Symmetry codes: (i)  $\frac{1}{2} - x, \frac{1}{2} - y, -z$ ; (ii)  $x, -y, z$ ; (iii)  $1 - x, y, -z$ ; (iv)  $\frac{1}{2} + x, \frac{1}{2} - y, z$ .]

Hence, (I) can be formulated as  $[\text{H}_{13}\text{O}_6]\{\text{trans}[\text{CrBr}_2(\text{Me}_2\text{tn})_2]\}_2\text{Br}_2(\text{ClO}_4)$ . For help in visualizing the complete structure of this compound, a drawing showing the  $[\text{H}_{13}\text{O}_6]^+$  cluster cation and the two conformational isomers of the  $\text{Cr}^{\text{III}}$  complex cation, together with two  $\text{Br}^-$  anions and a  $(\text{ClO}_4)^-$  anion, is included as Fig. 2.

## Experimental

The starting material *trans*- $[\text{CrCl}_2(\text{Me}_2\text{tn})_2]\text{ClO}_4$  was synthesized according to the literature method of House (1986). The complex *trans*- $[\text{CrCl}_2(\text{Me}_2\text{tn})_2]\text{ClO}_4$  (1.0 g) and NaOH (1.0 g) were suspended in water (10 ml) and the suspension heated at 333 K for 15 min. HBr (48%, 10 ml) and  $\text{HClO}_4$  (60%, 5 ml) were added to the mixture. The solution was concentrated at 333 K until green crystals appeared. The reaction mixture was then placed in an ice bath and the crystalline product which was deposited was filtered off, washed with cold 2-propanol and then with diethyl ether. Recrystallization of the crude product from 0.5 M HBr solution afforded green crystals of (I) suitable for X-ray analysis.

### Crystal data

$[\text{CrBr}_2(\text{C}_5\text{H}_{14}\text{N}_2)_2]_2\text{Br}_2 \cdot$ $\text{HClO}_4 \cdot 6\text{H}_2\text{O}$	$D_x = 1.731 \text{ Mg m}^{-3}$
$M_r = 1200.74$	Mo $K\alpha$ radiation
Monoclinic, $C2/m$	Cell parameters from 24 876 reflections
$a = 22.3862(8) \text{ \AA}$	$\theta = 1.6\text{--}27.5^\circ$
$b = 24.7393(8) \text{ \AA}$	$\mu = 5.78 \text{ mm}^{-1}$
$c = 8.3230(3) \text{ \AA}$	$T = 293(2) \text{ K}$
$\beta = 91.624(1)^\circ$	Plate, green
$V = 4607.6(3) \text{ \AA}^3$	$0.32 \times 0.32 \times 0.10 \text{ mm}$
$Z = 4$	

### Data collection

Rigaku R-Axis RAPID Imaging Plate diffractometer	5320 independent reflections
$\omega$ scans	4188 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (Higashi, 1995)	$R_{\text{int}} = 0.052$
$T_{\text{min}} = 0.188, T_{\text{max}} = 0.560$	$\theta_{\text{max}} = 27.5^\circ$
20 618 measured reflections	$h = -26 \rightarrow 26$
	$k = -30 \rightarrow 32$
	$l = -10 \rightarrow 10$

### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0520P)^2 + 9.0033P]$
$R[F^2 > 2\sigma(F^2)] = 0.042$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.108$	$(\Delta/\sigma)_{\text{max}} = 0.002$
$S = 1.05$	$\Delta\rho_{\text{max}} = 0.73 \text{ e \AA}^{-3}$
5320 reflections	$\Delta\rho_{\text{min}} = -0.60 \text{ e \AA}^{-3}$
228 parameters	
H-atom parameters constrained	

**Table 1**

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

Br11—Cr1	2.4876 (4)	Cr1—N11	2.084 (3)
Br21—Cr2	2.5017 (9)	Cr2—N21	2.078 (3)
Br22—Cr2	2.4495 (9)	Cr2—N22	2.089 (3)
Cr1—N12	2.078 (3)		
N12—Cr1—N11	91.46 (12)	N22—Cr2—N22 <sup>i</sup>	88.94 (18)
N12—Cr1—Br11	90.59 (9)	N21—Cr2—Br22	91.47 (9)
N11—Cr1—Br11	90.24 (9)	N22—Cr2—Br22	91.20 (10)
N21—Cr2—N21 <sup>i</sup>	89.79 (18)	N21—Cr2—Br21	88.45 (9)
N21—Cr2—N22	90.57 (13)	N22—Cr2—Br21	88.87 (10)
N21—Cr2—N22 <sup>i</sup>	177.29 (14)		

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

**Table 2**

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

$D\text{---}H\text{---}A$	$D\text{---}H$	$H\text{---}A$	$D\text{---}A$	$D\text{---}H\text{---}A$
N11—H11M $\cdots$ O2 <sup>i</sup>	0.90	2.57	3.438 (6)	163
N11—H11N $\cdots$ Br3 <sup>ii</sup>	0.90	2.77	3.661 (3)	169
N12—H12M $\cdots$ Br3	0.90	2.62	3.496 (3)	163
N12—H12N $\cdots$ Br3 <sup>iii</sup>	0.90	3.06	3.567 (3)	117
N21—H21M $\cdots$ Br3 <sup>iv</sup>	0.90	2.58	3.475 (3)	177
N21—H21N $\cdots$ Br11 <sup>v</sup>	0.90	2.90	3.580 (3)	134
N22—H22N $\cdots$ Br3 <sup>iv</sup>	0.90	2.62	3.516 (3)	176
O1—H00 $\cdots$ O1 <sup>vi</sup>	1.17	1.17	2.347 (9)	180
O1—H01A $\cdots$ O3 <sup>vii</sup>	0.90	1.77	2.667 (7)	175
O1—H01B $\cdots$ O2	0.90	1.82	2.662 (7)	155
O2—H02A $\cdots$ O42 <sup>iii</sup>	0.90	2.05	2.89 (1)	157
O2—H02B $\cdots$ Br3	0.90	2.61	3.420 (5)	150
O3—H03A $\cdots$ Br3 <sup>iii</sup>	0.90	2.70	3.383 (5)	133
O3—H03A $\cdots$ Br22 <sup>iii</sup>	0.90	3.02	3.575 (5)	121

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

The water H atoms were located from difference syntheses and refined as parts of rigid groups, with  $\text{O—H} = 0.90 \text{ \AA}$  and  $U_{\text{iso}}(\text{H}) = 0.044 \text{ \AA}^2$ . The position of the  $\text{H}^+$  ion (H00) was fixed at the midpoint between atoms O1 and O1( $1 - x, y, -z$ ). Other H atoms were placed geometrically and refined using a riding model, with  $\text{N—H} = 0.90 \text{ \AA}$ ,  $\text{C—H} = 0.96$  (for methyl H) or  $0.97 \text{ \AA}$  (for methylene H), and  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N or C})$ . The H atoms of the methyl groups (C24, C25, C29 and C30) were treated as positionally disordered over two sites.

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *TEXSAN* (Molecular Structure Corporation & Rigaku, 2000); program(s) used to solve

structure: *SHELXS86* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP* (Johnson, 1970); software used to prepare material for publication: *SHELXL97*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: OB1082). Services for accessing these data are described at the back of the journal.

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