

Pentaamminenitrosylchromium(III) dichloride

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Key indicators

Single-crystal X-ray study

T = 273 K

Mean $\sigma(\text{O}-\text{N}) = 0.008 \text{ \AA}$

Disorder in main residue

R factor = 0.044

wR factor = 0.117

Data-to-parameter ratio = 17.7

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

The cation in the title compound, $[\text{Cr}(\text{NO})(\text{NH}_3)_5]\text{Cl}_2$, is distorted octahedral with a linear Cr–N–O moiety. The short Cr–N(nitrosyl) distance of 1.692 (7) Å indicates significant multiple bonding between these atoms. The Cr atom, the NO group and the N atom *trans* to NO have *m2m* symmetry and the remaining N and Cl atoms have *m* symmetry.

Comment

This report is the first example of an X-ray structural study of a pentaamminenitrosylchromium complex, $[\text{Cr}(\text{NO})(\text{NH}_3)_5]^{2+}$, even though chromium complexes with Cl, Br, NO_3 , ClO_4 , $\text{B}(\text{C}_6\text{H}_5)_4$ and $\text{OC}_6\text{H}_2(\text{NO}_2)_3$ ligands have been characterized (Mori *et al.*, 1963; Griffith, 1963; Kobayashi *et al.*, 1969). The structure of the related compound $[\text{Cr}(\text{NO})(\text{H}_2\text{O})_5]\text{SO}_4$ (Ardon & Cohen, 1993) has been reported. The results of *ab initio* calculations on $[\text{Cr}(\text{NO})(\text{H}_2\text{O})_5]^{2+}$ have also been reported (Shim *et al.*, 1995).

The title compound, $[\text{Cr}(\text{NO})(\text{NH}_3)_5]\text{Cl}_2$ (Fig. 1 and Table 1), (I), was found to crystallize in the orthorhombic space group *Cmcm* with the Cr atom, the NO group and the N2 atom, *trans* to NO, having *m2m* symmetry imposing disorder in the N2H₃ ammine group. The remaining N atoms and the Cl atom are at sites with *m* symmetry.

The Cr^{III} centre exists in a slightly distorted octahedral geometry with a strictly linear Cr–N–O arrangement (from symmetry). The N(nitrosyl)–Cr–N(ammonia, *cis* to NO) angles are 94.2 (1) and 92.8 (1)° so that the Cr–N(ammonia, *cis* to NO) bonds are slightly bent towards the ammonia molecule in the *trans* position to NO.

The short Cr–N(nitrosyl) distance of 1.692 (7) Å indicates the presence of multiple bonding between these atoms. A short Cr–N(nitrosyl) distance of 1.682 (2) Å was also reported for $[\text{Cr}(\text{NO})(\text{H}_2\text{O})_5]^{2+}$ (Ardon & Cohen, 1993). The Cr–N(ammonia) distances in $[\text{Cr}(\text{NH}_3)_6]^{3+}$ are 2.070 (8) Å (Clegg, 1982) and 2.085 (10) Å (average; Dieterich & Strähle, 1993). These are close to those of Cr–N(ammonia, *cis* to NO) in $[\text{Cr}(\text{NO})(\text{NH}_3)_5]^{2+}$.

The O(nitrosyl) atom appears to form hydrogen bonds with both H3 and H4 of adjacent ammonia molecules, with distances and angles as shown in Table 2.

Experimental

The preparation of $[\text{Cr}(\text{NO})(\text{NH}_3)_5]\text{Cl}_2$ was carried out according to the literature method (Mori *et al.*, 1963). Crystals of $[\text{Cr}(\text{NO})(\text{NH}_3)_5]\text{Cl}_2$ were obtained by recrystallization from aqueous hydrochloric acid.

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Crystal data

 $[\text{Cr}(\text{NO})(\text{NH}_3)_5]\text{Cl}_2$
 $M_r = 238.06$

 Orthorhombic, *Cmcm*
 $a = 10.0236$ (9) Å

 $b = 9.098$ (3) Å

 $c = 10.357$ (1) Å

 $V = 944.5$ (5) Å³
 $Z = 4$
 $D_x = 1.674$ Mg m⁻³
 $D_m = 1.62$ Mg m⁻³
 D_m measured by flotation

 Mo $K\alpha$ radiation

Cell parameters from 1131

reflections

 $\theta = 3.0$ – 27.5°
 $\mu = 1.73$ mm⁻¹
 $T = 273$ K

Prism, orange

 $0.20 \times 0.10 \times 0.10$ mm

Data collection

Rigaku/MSM Mercury CCD

diffractometer

 ω scans

 Absorption correction: multi-scan
(*TEXSAN*; Molecular Structure
Corporation & Rigaku Corpora-
tion, 1999)

 $T_{\min} = 0.587$, $T_{\max} = 0.707$

5457 measured reflections

601 independent reflections

 389 reflections with $F^2 > 2\sigma(F^2)$
 $R_{\text{int}} = 0.084$
 $\theta_{\text{max}} = 27.2^\circ$
 $h = -11 \rightarrow 12$
 $k = -11 \rightarrow 11$
 $l = -13 \rightarrow 11$

Refinement

 Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.117$
 $S = 0.83$

601 reflections

34 parameters

H-atom parameters not refined

 $w = 1/[\sigma^2(F_o^2) + (0.034P^2)]$

 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.88$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.48$ e Å⁻³
Table 1

Selected geometric parameters (Å, °).

Cr1–N1	1.697 (7)	Cr1–N4	2.087 (4)
Cr1–N2	2.126 (7)	Cr1–N4 ⁱⁱ	2.087 (4)
Cr1–N3	2.085 (4)	O1–N1	1.179 (8)
Cr1–N3 ⁱ	2.085 (4)		
N1–Cr1–N2	180.0	N3–Cr1–N3 ⁱ	171.4 (3)
N1–Cr1–N3	94.3 (1)	N3–Cr1–N4	89.79 (1)
N1–Cr1–N4	92.9 (1)	N4–Cr1–N4 ⁱⁱ	174.3 (3)
N2–Cr1–N3	85.7 (1)	Cr1–N1–O1	180.0
N2–Cr1–N4	87.1 (1)		

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

Table 2

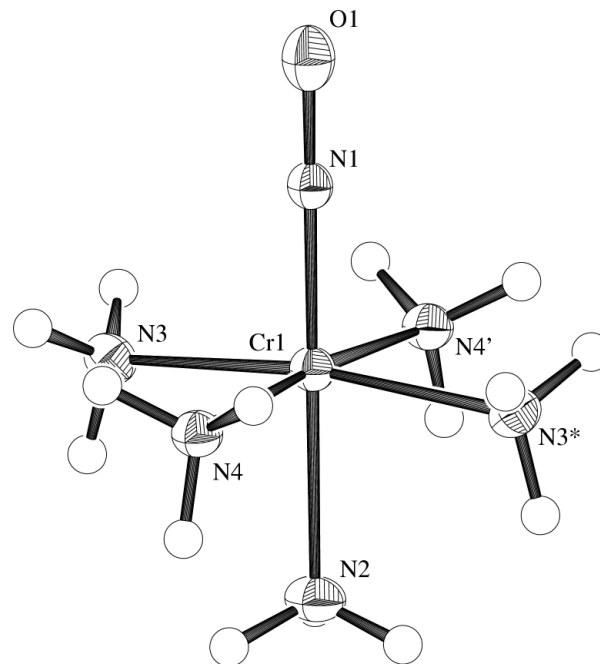
Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
N3–H3 [·] ··O1 ⁱ	0.95	2.75	3.302 (6)	118
N3 ⁱⁱ –H3 ⁱⁱ ··O1 ⁱⁱⁱ	0.95	2.75	3.302 (6)	118
N4–H6 [·] ··O1 ^{iv}	0.95	2.89	3.211 (5)	101
N4–H6 ⁱⁱ ··O1 ^{iv}	0.95	2.89	3.211 (5)	101
N4 ^v –H6 ^v ··O1 ^{vi}	0.95	2.89	3.211 (5)	101
N4 ^v –H6 ^{vii} ··O1 ^{vi}	0.95	2.89	3.211 (5)	101

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

All the H atoms were fixed in the refinement with isotropic displacement parameters set at 1.2 times the value of the equivalent isotropic displacement parameter of their carrier atom. Since the N2 atom is at a site with $m2m$ symmetry, the six disordered H atoms of the ammonia molecule have been developed by mirror symmetry.

Data collection: *CRYSTALCLEAR* (Rigaku, 1996); cell refinement: *CRYSTALCLEAR*; data reduction: *TEXSAN* (Molecular Structure Corporation & Rigaku Corporation, 1999); program(s)


Figure 1

The structure of the $[\text{Cr}(\text{NO})(\text{NH}_3)_5]^{2+}$ cation. Displacement ellipsoids are plotted at the 50% probability level (Johnson, 1976). For clarity, only three out of the six disordered H atoms of the ammonia molecule *trans* to NO developed by mirror symmetry are shown.

used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *TEXSAN*; molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *TEXSAN*.

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